PCT

WORLD INTELLECTUAL PROPERTY ORGANIZATION International Bureau



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification 6:

. . | (11) 11

(11) International Publication Number:

WO 99/45098

¢.

C12N 1/20, 15/74, A61K 48/00

A2

(43) International Publication Date:

10 September 1999 (10.09.99)

(21) International Application Number:

PCT/IB99/00587

(22) International Filing Date:

3 March 1999 (03.03.99)

(81) Designated States: AU, CA, CN, JP, KR, NZ, European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE).

(30) Priority Data:

09/036,582

6 March 1998 (06.03.98)

US

Published

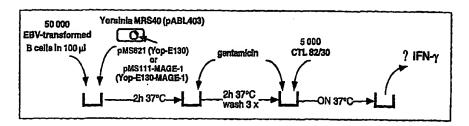
Without international search report and to be republished upon receipt of that report.

(71)(72) Applicants and Inventors: VAN DER BRUGGEN, Pierre, B. [BE/BE]; 12/2, chemin du Relai, B-1325 Corroy-Le-Grand (BE). CORNELIS, Guy, R. [BE/BE]; 2B, avenue des Anciens Combattants, B-1950 Kraainem (BE). BOLAND, Anne, M. [BE/BE]; 159, rue Theodore Decuyper, Boîte postale 043, B-1200 Bruxelles (BE). BOON-FALLEUR, Thierry, R. [BE/BE]; 12, rue du Buisson, B-1050 Bruxelles (BE).

(74) Agent: KOLB, Helga; Hoffmann, Eitle, Arabellastrasse 4, D-81925 München (DE).

(54) Title: DELIVERY OF PROTEINS INTO EUKARYOTIC CELLS WITH RECOMBINANT YERSINIA

Anti-MAGE-1.A1 CTL recognize HLA-A1 cells incubated with Yersinia which produces a YopE130.MAGE-1 fusion protein



(57) Abstract

The present invention relates to recombinant Yersinia and the use thereof for delivery of proteins into eukaryotic cells, including related compositions and methods of treatment and related assays.

FOR THE PURPOSES OF INFORMATION ONLY

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AL	Albania	ES	Spain	LS	Lesotho	SI	Slovenia
AM	Armenia	FI	Finland	LT	Lithuania	SK	Slovakia
AT	Austria	FR	France	LU	Luxembourg	SN	Senegal
AU	Australia	GA	Gabon	LV	Latvia	SZ	Swaziland
AZ	Azerbaijan	GB	United Kingdom	MC	Monaco	TD	Chad
BA	Bosnia and Herzegovina	GE	Georgia	MD	Republic of Moldova	TG	Togo
BB	Barbados	GH	Ghana	MG	Madagascar	TJ	Tajikistan
BE	Belgium	GN	Guinea	MK	The former Yugoslav	TM	Turkmenistan
BF	Burkina Faso	GR	Greece		Republic of Macedonia	TR	Turkey
BG	Bulgaria	HU	Hungary	ML	Mali	TT	Trinidad and Tobago
BJ	Benin	ΙE	Ireland	MN	Mongolia	UA	Ukraine
BR	Brazil	IL	Israel	MR	Mauritania	UG	Uganda
BY	Belarus	IS	Iceland	MW	Malawi	US	United States of America
CA	Canada	IT	Italy	MX	Mexico	UZ	Uzbekistan
CF	Central African Republic	JP	Japan	NE	Niger	VN	Viet Nam
CG	Congo	KE	Kenya	NL	Netherlands	YU	Yugoslavia
CH	Switzerland	KG	Kyrgyzstan	NO	Norway	zw	Zimbabwe
CI	Côte d'Ivoire	KP	Democratic People's	NZ	New Zealand		
CM	Cameroon		Republic of Korea	PL	Poland		
CN	China	KR	Republic of Korea	PT	Portugal		
CU	Cuba	KZ	Kazakstan	RO	Romania		
CZ	Czech Republic	LC	Saint Lucia	RU	Russian Federation		
DE	Germany	LI	Liechtenstein	SD	Sudan		
DK	Denmark	LK	Sri Lanka	SE	Sweden		
EE	Estonia	LR	Liberia	SG	Singapore		

DELIVERY OF PROTEINS INTO EUKARYOTIC CELLS WITH RECOMBINANT YERSINIA

The present invention relates to recombinant Yersinia and the use thereof for delivery of proteins into eukaryotic cells.

5

10

15

20

25

30

Bacteria of the genus Yersinia cause diseases in humans and rodents ranging from enteritis and lymphadenitis to plague. The genus Yersinia encompasses three species: Yersinia enterocolitica, which is the most prevalent Yersinia species in humans and causes a broad range of gastro-intestinal syndromes; Yersinia pseudotuberculosis, which causes adenitis and septicaemia; and Yersinia pestis, which is the causative agent of plague.

In spite of the differences in the infection routes, these three species of Yersinia share a common capacity to resist the non-specific immune response of the human or rodent host and to proliferate in the host lymphatic tissues. Anatomo-pathological examinations revealed that Yersinia are not detected inside the inflammatory or parenchymal cells of the infected animals (Simonet et al. (1990) Infect. Immun. 58: 841-845). Consistent with these in vivo observations, Yersinia are resistant to phagocytosis in vitro by macrophages and polymorphonuclear leukocytes. See review by Cornelis et al. (1997) Mol. Microbiol. 23(5): 861-867. Yersinia enterocolitica also has the capacity to enter certain cultured epithelial cells, a process generally referred to as

invasion (Miller et al. (1988) *Infect. Immun.* 56: 1242-1248).

Genetic studies revealed that the virulence of Yersinia is determined by a 70 kb plasmid (pYV), 5 which encodes and governs the production of a set of proteins called Yops (for Yersinia outer proteins). These Yops form an integrated anti-host system that allows the extracellular adhesion of Yersinia to the surface of host cells and the subsequent injection of a set of toxic effector proteins into the host cell's 10 cytosol. Recent studies further revealed that such an anti-host system, also called "Yersinia virulon", is composed of the following four elements: (i) a contact or type III secretion system called Ysc, which is devoted to the secretion of Yop proteins out of the 15 bacterial cells; (ii) a set of "translocators" for translocating the effector proteins into the eukaryotic host cells, which consist of YopB, YopD and possible other proteins such as LcrV; (iii) a control 20 element and recognition system (YopN and LcrG); and (iv) a set of "effector proteins" including YopE, YopH, YopO/YpkA, YopM and YopP/YopJ, which are injected (or translocated) into the eukaryotic host cells and disrupt the functions of such host cells. 25 Transcription of these genes is controlled both by temperature and by contact with a eukaryotic cell. See review by Cornelis et al. (1997).

The effector proteins disrupt the function of host cells in a number of ways. The 23 kd YopE is a cytotoxin that disrupts the actin-microfilament structure of cultured Hela cells (Rosqvist et al.

30

(1990) Mol. Microbiol. 4: 657-667; Rosqvist et al. (1991) Infect. Immun. 59: 4562-4569). The 51 kd YopH is a protein tyrosine phosphatase (PTPase) related to eukaryotic PTPases, which acts on tyrosine-5 phosphorylated proteins of infected macrophages (Hartland et al. (1994) Infect. Immun. 62: 4445-4453). Presumably as a result of this action, YopH inhibits bacterial uptake and oxidative burst by cultured macrophages (Rosquvist et al. (1988) Infect. Immun. 56: 2139-2143; Bliska et al. (1995) Infect. Immun. 63: 10 681-685). YopO (or YpkA) is an 81 kd serine/threonine kinase, which is targeted to the inner surface of the plasma membrane of the eukaryotic cell and might function to interfere with the signal transduction 15 pathway of the eukaryotic cell (Hakansson et al. (1996) Mol. Microbiol. 20: 593-603). YopM is an acidic 41 kd protein having 12 leucine-rich repeats, which suggests that YopM might bind thrombin and interfere with platelet-mediated events of the 20 inflammatory response (Leung et al. (1989) J. Bacterial. 171: 4623-4632). YopP is involved in the induction of apoptosis in macrophages (Mills et al. (1997) Proc. Acad. Natl. Sci. USA 94: 12638-12643). The molecular structures of these effector 25 proteins have been investigated to determine the elements in each effector protein that are required for their secretion and translocation. For this purpose, hybrid proteins have been engineered by

calmodulin-activated adenylate cyclase domain (or Cya)

fusing truncated Yop effector proteins of different length with certain reporter enzymes such as the

30

of the Bordetella pertussis cyclolysin. Successful secretion and/or translocation events could be detected by assays based on the enzymatic activity of the reporter protein. Sory et al. disclose, by applying this approach, that YopE and YopH of Y. enterocolitica are modular proteins composed of three domains, i.e., an N-terminal domain required for secretion, a translocation domain required for translocation into cells, and a C-terminal catalytic domain responsible for the toxic effector activity. Sory et al. (1995) Proc. Natl. Acad. Sci. USA 92: 11998-12002. The same domain organization has been demonstrated for YopM of Y. enterocolitca (Boland et al. (1996) EMBO J. 15: 5191-5201).

5

10

15

20

25

30

The present invention provides recombinant Yersinia for safe delivery of proteins into eukaryotic Such Yersinia are deficient in the production of functional effector proteins, but are endowed with a functional secretion and translocation system. present invention further provides expression vectors for use in combination with such mutant Yersinia for safe and efficient delivery of desired proteins into eukaryotic cells. This approach is useful not only for studying the function of a given protein, but also for designing therapeutic approaches. For example, a protein of a pathogenic origin, e.g., a tumor associated protein, a parasite antigen, or a viral antigen, can be delivered using the recombinant Yersinia of the present invention into antigen presenting cells for inducing an immune response specific for such a protein.

Most progressively growing neoplastic cells express potentially immunogenic tumor-associated antigens (TAAs), also called tumor rejection antigens (TRAs). TRAs, like other antigenic epitopes, are presented at the surface of tumor cells by MHC molecules and have been shown to induce a CTL response in vivo and in vitro. See, for example, van der Bruggen et al. (1991) Science 254: 1643-1647. However, such TRA-expressing tumor cells do not provoke reliable anti-tumor immune responses in vivo that are capable of controlling the growth of malignant cells. Boon et al. (1992) Cancer Surveys 23-37; T. Boon (1993) Int. J. Cancer 54: 180; T. Boon (1992) Advances Cancer Res. 58: 177-209. A number of genes have been identified that encode tumor rejection antigen precursors (or TRAPs), which are processed into TRAs in tumor cells. TRAP-encoding genes include members of the MAGE family, the BAGE family, the DAGE/Prame family, the GAGE family, the RAGE family, the SMAGE family, NAG, Tyrosinase, Melan-A/MART-1, gp100, MUC-1, TAG-72, CA125, mutated proto-oncogenes such as p21ras, mutated tumor suppressor genes such as p53, tumor associated viral antigens such as HPV16 E7. See, e.g., review by Van den Eynde and van der Bruggen (1997) in Curr. Opin. Immunol. 9:684-693, Sahin et al. (1997) in Curr. Opin. Immunol. 9:709-716, and Shawler et al. (1997) Advances in Pharmacology 40: 309-337 Academic Press, San Diego, California. The identification of

5

10

15

20

25

30

these genes has allowed recombinant production of TRAs

or TRAPs which may be subsequently used as vaccines to treat various cancerous conditions.

The present invention contemplates the use of recombinant Yersinia for delivery of desired proteins into eukaryotic cells. Particularly, the recombinant Yersinia of the present invention are useful for delivery of proteins or derivatives thereof to antigen presenting cells. In accordance with the present invention, antigen presenting cells upon receiving the delivery, present antigenic epitopes which can be recognized by T cells. Thus, the recombinant Yersinia of the present invention can be employed in a number of immune diagnostic or therapeutic approaches.

5

10

15

20

25

30

The present invention is further elaborated upon the following disclosure.

The present invention relates to recombinant *Yersinia* and the use thereof for delivery of proteins into eukaryotic cells.

One embodiment of the present invention provides mutant Yersinia strains deficient in producing functional effector proteins. A preferred mutant Yersinia strain of the present invention is a quintuple mutant strain designated as yopEHMOP.

Another embodiment of the present invention provides expression vectors for delivery of heterologous proteins to eukaryotic cells. In accordance with the present invention, such an expression vector is characterized by (in the 5' to 3' direction) a promoter, a first nucleic acid sequence encoding a delivery signal, a second nucleic acid

sequence fused thereto coding for a heterologous protein to be delivered.

5

10

15

20

25

30

According to this embodiment of the present invention, the promoter is preferably one from a Yersinia virulon gene; more preferably, an effectorencoding gene; even more preferably, a YopE gene. delivery signal is a polypeptide sequence from an Yersinia effector, including YopE, YopH, YopO/YpkA, YopM, and YopP/YopJ. Such delivery signal can be recognized by the secretion and translocation system of Yersinia. The heterologous protein of the present invention includes naturally occurring proteins or parts thereof such as tumor-associated proteins or known antigens of pathogens. The heterologous protein of the present invention also includes artificially designed proteins such as in-frame fusion of proteins or parts of proteins.

Yet another embodiment of the present invention provides recombinant Yersinia, i.e., yersinia of the above-described mutant strains further transformed with the expression vector of the present invention. Such recombinant Yersinia is preferred for delivery of heterologous proteins into eukaryotic cells. A preferred eukaryotic cell is an antigen presenting cell capable of presenting immunogenic epitopes derived from the heterologous proteins being delivered.

In a further aspect of the present invention, such recombinant *Yersinia* are contemplated in immunogenic compositions and methods for inducing an immune response, either a cellualr immune response

or a humoral immune response, or a combination of both.

5

10

15

20

25

30

Further to this aspect of the invention, the recombinant Yersinia of the present invention can be employed in an in vitro regime for assessing the efficacy of a vaccination regimen. The recombinant Yersinia of the present invention can also be employed in an ex vivo regime to generate specific CTLs and to use such CTLs for treating various pathological conditions such as tumors or infections by pathogens. The recombinant Yersinia of the present invention can also be employed in an in vivo regime, i.e., as a recombinant vaccine, to treat subjects suffering pathological conditions such as tumors or infections by pathogens.

Figure 1 illustrates the plasmid map of the expression vector pMS111-MAGE-1 (YopE $_{130}$ -MAGE1).

Figure 2 (A) depicts the procedure for stimulating CTL 82/30 with EBV-transformed human B cells (HLA-A1) mixed wth recombinant Yersinia; (B) depicts the quantitation of IFN- γ released by activated CTLs.

Figure 3 depicts the sequence of the Yersinia enterocolitica YopM gene.

Figure 4 depicts the sequence of the Yersinia enterocolitica YopE gene.

Figure 5 depicts the sequence of the Yersinia enterocolitica YopH gene.

Figure 6 depicts the sequence of the Yersinia enterocolitica YopP gene.

Figure 7 depicts the sequence of the Yersinia enterocolitica YopP gene.

5

30

The present invention relates to recombinant Yersinia and the use thereof for delivery of proteins into eukaryotic cells.

In particular, the present invention provides mutant Yersinia strains that are deficient in producing functional effector proteins. The present invention further provides an expression 10 vector, which, when transformed into a Yersinia of the above-described mutant strains, permits delivery of a desired protein, e.g., a MAGE-1 protein, into a eukaryotic cell, for example, a EBV-transformed human B cell. The present invention provides that, upon 15 receiving the delivered protein, the antigen presenting cell processes the delivered protein and presents antigenic epitopes in the context of an MHC molecule leading to an immune response specific for the protein being delivered. For example, human B 20 cells after taking up the MAGE-1 protein from a recombinant Yersinia of the present invention are recognized by a CTL clone specific for a MAGE-1 epitope. Thus the present invention provides an effective recombinant Yersinia system with diminished 25 toxicity for delivery of desired proteins into eukaryotic cells.

The term "Yersinia" as used herein means all species of Yersinia, including Yersinia enterocolitica, Yersinia pseudotuberculosis and Yersinia pestis.

For the purpose of the present invention, the term "recombinant Yersinia" used herein refers to Yersinia genetically transformed with the expression vectors of the present invention.

5

10

15

20

25

30

The term "delivery" used herein refers to the transportation of a protein from a Yersinia to a eukaryotic cell, including the steps of expressing the protein in the Yersinia, secreting the expressed protein(s) from such Yersinia and translocating the secreted protein(s) by such Yersinia into the cytosol of the eukaryotic cell. Accordingly, a "delivery signal" refers to a polypeptide sequence which can be recognized by the secretion and translocation system of Yersinia and directs the delivery of a protein from Yersinia to eukaryotic cells.

As used herein, the "secretion" of a protein refers to the transportation of such protein outward across the cell membrane of a Yersinia. The "translocation" of a protein refers to the transportation of such protein across the plasma membrane of a eukaryotic cell into the cytosol of such eukaryotic cell.

"Eukaryotic cells" as used herein, the surface of which Yersinia adhere to, are also referred to as "target cells" or "target eukaryotic cells".

One embodiment of the present invention provides mutant *Yersinia* strains which are deficient in producing functional effector proteins.

The effector proteins of Yersinia, i.e., the Yersinia virulon proteins which are normally translocated into the cytosol of the target eukaryotic

cells, are toxic to the target cell. Thus, a "functional effector protein" refers to an effector protein having a defined catalytic activity and which is capable of eliciting specific toxicity toward the target cells.

5

10

15

20

25

30

Accordingly, the mutant Yersinia of the present invention are used for delivery of proteins to eukaryotic cells, with diminished toxicity, i.e., toxicity which does not completely disable or kill the target cell. For the purpose of delivering proteins, the secretion and translocation system of the instant mutant Yersinia need to be intact.

Five effector genes have been cloned from Y. enterocolitica which are YopE, YopH, YopO, YopM, and YopP (Figures 3-7). The equivalent effector genes have been cloned from Y. pseudotuberculosis and are named as YopE, YopH, YpkA, YopM, and YopJ, respectively. Some effector genes have also been cloned from Y. pestis. The nucleic acid sequences of these Yop genes are available to those skilled in the art, e.g., in the Genebank Database

For the purpose of the present invention, the effector-encoding genes are denoted by italicized letters to be distinguished from the effector proteins. Mutant effector genes are denoted by letters of lower case. For example, YopE refers to the effector protein encoded by the YopE gene. YopE represents the wild type gene, while yopE represents a gene having a mutation.

According to the present invention, a mutant Yersinia strain can be generated by introducing at

least one mutation into at least one effector-encoding gene. Preferably, such effector-encoding genes include YopE, YopH, YopO/YpkA, YopM, and YopP/YopJ. The skilled artisan may employ any number of standard techniques to generate mutations in these Yop genes. Sambrook et al. describe in general such techniques. See Sambrook et al. (1989) Molecular Cloning: A Laboratory Manual, Second Edition Cold Spring Harbor Laboratory Press: Cold Spring Harbor, New York.

5

10

15

20

25

30

The term "mutation" is used herein as a general term and includes changes of both single base pair and multiple base pairs. Such mutations may include substitutions, frame-shift mutations, deletions and truncations.

In accordance with the present invention, the mutation can be generated in the promoter region of an effector-encoding gene so that the expression of such effector gene is abolished.

The mutation can also be generated in the coding region of an effector-encoding gene such that the catalytic activity of the encoded effector protein is abolished. The "catalytic activity" of an effector protein refers to the anti-target cell function of an effector protein, i.e., toxicity. Such activity is governed by the catalytic motifs in the catalytic domain of an effector protein. The approaches for identifying the catalytic domain and/or the catalytic motifs of an effector protein are well within the ken of those skilled in the art. See, for example, Sory et al. (1995), Boland et al. (1996) and Cornelis et al. (1997).

Accordingly, one preferred mutation of the present invention is a deletion of the entire catalytic domain. Another preferred mutation is a frameshift mutation in an effector-encoding gene such that the catalytic domain is not present in the protein product expressed from such "frameshifted" gene. A most preferred mutation is a mutation with the deletion of the entire coding region.

5

10

15

20

25

30

Other mutations are also contemplated by the present invention, such as small deletions or base pair substitutions, which are generated in the catalytic motifs of an effector protein leading to destruction of the catalytic activity of a given effector protein.

The mutations that are generated in the Yop genes may be introduced into Yersinia by a number of methods. One such method involves cloning a mutated Yop gene (i.e., a yop gene), into a "suicide" vector which is capable of introducing the mutated yop sequence into Yersinia via allelic exchange. Such "suicide" vectors are described by Kaniga et al. (1991) Gene 109: 137-141 and by Sarker et al. (1997) Mol. Microbiol 23: 409-411. In this manner, mutations generated in multiple Yop genes may be introduced successively into Yersinia, giving rise to polymutant recombinant Yersinia. The order in which these mutated yop sequences are introduced is not important.

A preferred mutant Yersinia strain of the present invention is a quintuple-mutant Yersinia strain in which all the effector-encoding genes are

mutated such that the resulting Yersinia no longer produce any functional effector proteins. Such quintuple-mutant Yersinia strain is designated as yopEHOMP for Y. enterocolitica or yopEHAM for Y. pseudotuberculosis. One example of such yopEHOMP strain is Y. enterocolitica MRS40 (pABL403).

5

10

15

20

25

30

Under some circumstances, it may be desired to mutate only some but not all of the effector Yop genes. For example, when a delivery is intended to target a macrophage, YopH is preferably not mutated since YopH is understood to inhibit the phagocytosis by macrophages. Rosqvist et al. (1988) and Rosqvist et al. (1989). Accordingly, the present invention further contemplates polymutant Yersinia other than quintuple-mutant Yersinia, e.g., double-mutant, triple-mutant, and quadruple-mutant Yersinia.

Alternatively, the quintuple-mutant strain yopEHOMP may still be used for delivery to macrophages, in which case a wild type YopH gene can be introduced into the quintuple-mutant Y. yopEHOMP strain by various known transformation procedures which are further described hereinafter. In this manner, a polymutant Yersinia strain can be generated in which a desired set of Yop genes are mutated such that only the protein of interest is delivered into the target cells.

A further aspect of the present invention is directed to an expression vector for use in combination with the instant mutant *Yersinia* strains to deliver a desired protein into eukaryotic cells. In accordance with the present invention, such a

vector is characterized by (in the 5' to 3' direction) a promoter, a first nucleic acid sequence encoding a delivery signal, a second nucleic acid sequence fused thereto coding for a heterologous protein to be delivered.

5

10

15

20

25

30

In accordance with present invention, the promoter of the expression vector is preferably from a Yersinia virulon gene. A "Yersinia virulon gene" refers to genes on the Yersinia pYV plasmid, the expression of which is controlled both by temperature and by contact with a target cell. See review by Cornelis et al. (1997). Such genes include genes coding for elements of the secretion machinary (the Ysc genes), genes coding for translocators (YopB, YopD, and LcrV), genes coding for the control elements (YopN and LcrG), and genes coding for effectors (YopE, YopH, YopO/YpkA, YopM and YopP/YopJ).

In a preferred embodiment of the present invention, the promoter is from an effector-encoding gene selected from any one of YopE, YopH, YopO/YpkA, YopM and YopP/YopJ. More preferrably, the promoter is from YopE.

Further in accordance with the present invention, a first DNA sequence coding for a delivery signal is operably linked to the promoter.

"A delivery signal", as described hereinabove, refers to a polypeptide which can be recognized by the secretion and translocation system of *Yersinia* and therefore directs the secretion and translocation of a protein into a eukaryotic cell.

According to the present invention, such a polypeptide is from an effector protein. The effector proteins include YopE, YopH, YopO/YpkA, YopM, and YopP/YopJ. Preferably, the effector protein is YopE. More preferably, the effector protein is YopE of Yersinia enterocolitica.

5

10

15

20

25

30

One skilled in the art is familiar with the methods for identifying the polypeptide sequences of an effector protein that are capable of delivering a protein. For example, one such method is described by Sory et al. (1994). Briefly, polypeptide sequences from various portions of the Yop proteins can be fused in-frame to a reporter enzyme such as the calmodulinactivated adenylate cyclase domain (or Cya) of the Bordetella pertussis cyclolysin. Delivery of a Yop-Cya hybrid protein into the cytosol of eukaryotic cells is indicated by the appearance of cyclase activity in the infected eukaryotic cells that leads to the accumulation of cAMP. Examples of such delivery signal polypeptides include from Y. enterocolitica: YopE₁₃₀ (the N-terminal 130 amino acids of YopE), YopE₅₀, YopM₁₀₀ and YopH₇₁.

By employing such an approach, one skilled in the art can determine, if desired, the minimal sequence requirement, i.e., a contiguous amino acid sequence of the shortest length, that is capable of delivering a protein. See, e.g., Sory et al (1994). Accordingly, preferred delivery signals of the present invention consists of at least the minimal sequence of amino acids of a Yop effector protein that is capale of delivering a protein.

Further in accordance with the present invention, a second DNA sequence encoding a heterologous protein is fused in frame to the first DNA sequence in the instant vector for delivery into eukaryotic cells.

5

10

15

20

25

30

The term "heterologous protein" used herein refers to a protein other than a Yersinia Yop protein. "Yop proteins" refer to Yersinia virulon proteins that are secreted, including the translocators and the effectors.

According to the present invention, "a heterologous protein" includes naturally occuring proteins or parts thereof. The term "part of a protein" includes a peptide or polypeptide fragment of a protein that is of sufficient length to be antigenic. Preferably, such a fragment consists of at least 8 or 9 contiguous amino acids of a protein. "A heterologous protein" as used in the present invention also includes artificially engineered proteins or parts thereof, such as fusion of two or more naturally occurring proteins or parts thereof, polyepitopes (inframe fusion of two or more peptide epitopes) as exemplified by Thompson et al. (1995) in *Proc. Natl. Acad. Sci. USA* 92: 5845-5849.

The protein expressed from the fused first and second DNA sequences is also termed as a "fusion protein" or a "hybrid protein", i.e., a hybrid of *Yersinia* delivery signal and a heterologous protein.

There is no particular limitation in the heterologous protein that can be delivered. The present invention particularly contemplates proteins,

such as, e.g., known tumor associated proteins, known antigens of pathogens, and cytokines.

5

10

15

20

25

30

.A "tumor associated protein" refers to a protein that is specifically expressed in tumors or expressed at an abnormal level in tumors relative to normal tissues. Such tumor associated proteins include, but are not limited to, members of the MAGE family, the BAGE family (such as BAGE-1), the DAGE/Prame family (such as DAGE-1), the GAGE family, the RAGE family (such as RAGE-1), the SMAGE family, NAG, Tyrosinase, Melan-A/ MART-1, gp100, MUC-1, TAG-72, CA125, mutated proto-oncogenes such as p21ras, mutated tumor suppressor genes such as p53, tumor associated viral antigens (e.g., HPV16 E7), HOM-MEL-40, HOM-MEL-55, NY-COL-2, HOM-HD-397, HOM-RCC-1.14, HOM-HD-21, HOM-NSCLC-11, HOM-MEL-2.4, and HOM-TES-11. Members of the MAGE family include, but are not limited to, MAGE-1, MAGE-2, MAGE-11. Members of the GAGE family include, but are not limited to, GAGE-1, GAGE-6. See, e.g., review by Van den Eynde and van der Bruggen (1997) in Curr. Opin. Immunol. 9:684-693, Sahin et al. (1997) in Curr. Opin. Immunol. 9:709-716, and Shawler et al. (1997). These proteins have been shown to associate with certain tumors such as melanoma, lung cancer, prostate cancer, breast cancer, renal cancer and others.

A number of known antigens derived from pathogens can also be employed according to the present invention. Pathogens contemplated by the present invention include viruses, bacteria, parasites and fungi. Specific examples of antigens

characteristic of a pathogen include the influenza virus nucleoprotein (residues 218-226, as set forth in F. et al. (1997) J. Virol. 71: 2715-2721) antigens from Sendai virus and lymphocytic choriomeningitis 5 virus (see, An et al. (1997) J. Virol. 71: 2292-2302), the B1 protein of hepatitis C virus (Bruna-Romero et al. (1997) Hepatology 25: 470-477), the virus envelope glycoprotein gp 160 of HIV (Achour et al. (1996) J. Virol. 70: 6741-6750), amino acids 252-10 260 or the circumsporozite protein of Plasmodium berghei (Allsopp et al. (1996) Eur. J. Immunol. 26: 1951-1958), the influenza A virus nucleoprotein (residues 366-374, Nomura et al. (1996) J. Immunol. Methods 193: 4149), the listeriolysin O protein of 15 Listeria monocytogenes (residues 91-99, An et al. (1996) Infect. Immun. 64: 1685-1693), the E6 protein (residues 131-140, Gao et al. (1995) J. Immunol. 155: 5519-5526) and E7 protein (residues 21-28 and 48-55, Bauer et al. (1995) Scand. J. Immunol. 42: 317-323) 20 of human papillomavirus type 16, the M2 protein of respiratory syncytial virus (residues 82-90 and 81-95, Hsu et al. (1995) Immunology 85: 347-350), the herpes simplex virus type 1 ribonucleotide reductase (see, Salvucci et al. (1995) J. Gen. Virol. 69: 1122-1131) 25 and the rotavirus VP7 protein (see, Franco et al. (1993) J. Gen. Virol. 74: 2579-2586), P. falciparum antigens (causing malaria) and hepatitis B surface antigen (Gilbert et al. (1997) Nature Biotech. 15: 1280-1283).

Accordingly, sequences coding for the abovedescribed proteins may be cloned into the present expression vector for delivery.

5

10

15

20

25

30

A number of coding sequences for small antigenic peptides can also be employed in the present invention. One skilled in the art can readily determine the length of the fragments required to produce immunogenic peptides. Alternatively, the skilled artisan can also use coding sequences for peptides that are known to elicit specific T cell responses, such as tumor-associated antigenic peptides (TRAs) as disclosed by U.S. Patent No. 5,462,871, U.S. Patent No. 5,558,995, U.S. Patent No. 5,554,724, U.S. Patent No. 5,585,461, U.S. Patent No. 5,591,430, U.S. Patent No. 5,554,506, U.S. Patent No. 5,487,974, U.S. Patent No. 5,530,096, U.S. Patent No. 5,519,117. Examples of TRAs are provided in Table 1. See also review by Van den Eynde and van der Bruggen (1997) and Shawler et al. (1997). Antigenic peptides of a pathogen origin can also be used, such as those

As described herein above, sequences coding for a full-length naturally occurring protein, a part of a naturally occurring protein, combinations of parts of a naturally occurring protein, or combinations of different naturally occurring proteins or parts from different proteins, may all be employed in the present invention. For example, a sequence coding for multiple epitopes may be used, such as those described by Thomson et al. (1995). Preferably, the second DNA sequence of the present expression

disclosed by Gilbert et al. (1997).

vector codes for at least one epitope of a protein. An "epitope" refers to a peptide of at least 8 or 9 amino acids.

5

10

Those skilled in the art are familiar with the techniques to make DNA fragments coding for a part of a protein, or link a DNA sequence encoding a part of one protein in frame to a DNA sequence encoding a part of another protein and the like.

The vectors of the instant invention may include other sequence elements such as a 3' termination sequence (including a stop codon and a poly A sequence), or a gene conferring a drug resistance which allows the selection of Yersinia transformants having received the instant vector.

15 The expression vectors of the present invention may be transformed by a number of known methods into Yersinia. For the purpose of the present invention, the methods of transformation for introducing an expression vector include, but are not 20 limited to, electroporation, calcium phosphate mediated transformation, conjugation, or combinations thereof. For example, a vector can be transformed into a first bacteria strain by a standard electroporation procedure. Subsequently, such a 25 vector can be transferred from the first bacteria strain into Yersinia by conjugation, a process also called "mobilization". Yersinia transformant (i.e., Yersinia having taken up the vector) may be selected, e.g., with antibiotics. These techniques are well 30 known in the art. See, for example, Sory et al.

known in the art. See, for example, Sory et al (1994).

One preferred embodiment of the present invention is directed to a *Yersinia* of the above-described mutant *Yersinia* strain transformed with an expression vector for delivery of a heterologous protein as herinabove described into a eukaryotic cell.

Accordingly, the present invention contemplates a method for delivering heterologous proteins as hereinabove described into eukaryotic cells.

5

10

15

20

25

The present invention contemplates a wide range of eukaryotic cells that may be targeted by the instant recombinant Yersinia.

By "target", is meant the extracellular adhesion of Yersinia to a eukaryotic cell.

In particular, the present invention contemplates antigen-presenting cells. "Antigen presenting cells" as referred herein express at least one class I or class II MHC determinant and may include those cells which are known as professional antigen-presenting cells such as macrophages, dendritic cells and B cells. Other professional antigen-presenting cells include monocytes, marginal zone Kupffer cells, microglia, Langerhans' cells, interdigitating dendritic cells, follicular dendritic cells, and T cells. Facultative antigen-presenting cells can also be used according to the present invention. Examples of facultative antigen-presenting cells include astrocytes, follicular cells,

30 endothelium and fibroblasts. As used herein,

"antigen-presenting cells" encompass both professional and facultative types of antigen-presenting cells.

The antigen presenting cells can be isolated from tissue or blood (containing peripheral blood mononuclear cells) samples obtained from a mammal such as a human or rodent. Cell lines established from such samples may also be used. Procedures for establishing cell lines are well known in the art. Certain cell lines may be obtained directly from the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, 20852-1776. Both normal and malignant cells may be employed.

5

10

15

20

25

30

In accordance with a preferred embodiment of the present invention, the MHC determinants expressed by the antigen presenting cell are compatible with those expressed by the mammal involved, and at least one of these MHC determinants is capable of presenting one or more antigenic epitopes derived from the protein being delivered.

One skilled in the art is also familiar with the methods for determining whether the MHC molecules expressed by the antigen presenting cell are compatible with those of the mammal subject involved, such as well known HLA-typing procedures. See general teachings by Coligan et al. (1994) Current Protocols in Immunology John Wiley & Sons Inc: New York, New York.

Those skilled in the art are able, through the extensive teachings in the art, to determine the MHC molecule for presentation of a particular antigen. For example, U.S. Patent No. 5,405,940 teaches the

determination of HLA-Al as the presenting molecule for a peptide of MAGE-1, EADPTGHSY; U.S. Patent No. 5,558,995 teaches the determination of HLA-Cw1601 for presenting another peptide of MAGE-1, SAYGEPRKL; U.S. Patent No. 5,530,096 teaches the determination of HLA-A2 as the presenting molecule for a peptide of Tyrosinase, MLLAVLYCL. In the event the eukaryotic cells being targeted do not express a desired HLA or MHC molecule, the gene encoding such molecule may be introduced into the eukaryotic cells by well known transformation or transfection procedures.

5

10

15

20

25

30

Further in accordance with the present invention, the delivery of a protein can be achieved by contacting a eukaryotic cell with a recombinant Yersinia under appropriate conditions. Various references and techniques are conventionally available for those skilled in the art regarding the conditions for inducing the expression and translocation of virulon genes, including the desired temperature, Ca++ concentration, manners in which Yersinia and target cells are mixed, and the like. See, for example, Cornelis, Cross talk between Yersinia and eukaryotic cells, Society for General Microbiology Symposium, 55; MoCRAE, SAUNDERS, SMYTH, STOW (eds), Molecular aspects of host-pathoge interactions, Cambridge University Press, 1997. The conditions may vary depending on the type of eukaryotic cells to be targeted, e.g., the conditions for targeting human epithelial carcinoma Hela cells (Sory et al. (1994)); the conditions for targeting mouse thymoma or melanoma cells (Starnbach et al. (1994) J. Immunol. 153: 1603); the conditions

for targeting mouse macrophages (Boland et al. (1996)). Such variations can be addressed by those skilled in the art using conventional techniques.

Those skilled in the art can also use a 5 number of assays to determine whether the delivery of a fusion protein is successful. For example, the fusion protein may be labeled with an isotope or an immunofluoresceine, or detected by a immunofluoresceine conjugated antibody, as disclosed 10 by Rosqvist et al. (1994) EMBO J. 13: 964. determination can also be based on the enzymatic activity of the protein being delivered, e.g., the assay described by Sory et al. (1994). The determination can also be based on the antigenicity of 15 the protein being delivered. For example, the delivery of a MAGE-1 protein into EBV-transformed human B cells can be detected by the recognition of such targeted B cells by CTL cells specific for MAGE-1 epitopes. Such CTL recognition, in turn, may be 20 detected by a number of assays including assaying the secretion of IFN-y from the activated CTLs or Cr51 release from lysed target cells. Methods such as Western-blot analysis using antibodies specific against the protein being delivered, PCR in situ 25 hybridization, or ELISPOT (Mabtech AB, Sweden) may also be employed for such determination. See, e.g., W. Herr et al. (1997) J. Immunol. Methods 203: 141-152 and W. Herr et al. (1996) J. Immunol. Methods 191:

In a further aspect of the present invention, recombinant *Yersinia* capable of delivering

131-142.

proteins to antigen-presenting cells are employed for inducing an immune response. Accordingly, the present invention contemplates immunogenic compositions and methods for inducing specific immune responses using the instant recombinant Yersinia as described hereinabove.

The immune responses contemplated by the present invention include cellular immune responses (mediated primarily by T cells) and humoral immune responses (mediated primarily by antibodies). Janeway and Travers teach in general these immune response. (Janeway and Travers (1996) Immunology, The Immune System in Health and Disease 2nd ed. Garland Publishing, Inc.: New York, New York, and London, England.) (See also, review by O. Tureci et al. (1997) Molecular Medicine Today 3(8): 342-349.

According to this aspect of the present invention, the immune responses induced with recombinant Yersinia can be utilized in a number of regimes for diagnostic or therapeutic use. For example, recombinant Yersinia can be employed in an in vitro procedure for monitoring the efficacy of a vaccination therapy in a mammal such as a human or rodent. In this regime, certain antigen presenting cells (e.g., dendritic cells) are taken from a subject being vaccinated with immunogenic compositions, e.g., a particular antigen. Such antigen presenting cells are then contacted with recombinant Yersinia capable of delivering the antigen which is used for vaccination. Subsequently, peripheral blood lymphocytes taken from the same subject (i.e.,

autologous PBLs) are added, preferably in combination with cytokines such as IL-2, to the mixture of antigen presenting cells and Yersinia. The efficacy of the vaccination can be assessed after priming and then after successive boosts by the presence of CTLs or antibodies that are specific for the relevant antigen. The presence of specific CTLs can be detected using standard assays such as an assay for Cr⁵¹ release or for the secretion of IFN-gamma. The presence of specific antibodies can be detected by assays such as ELISA using the antigens which are immobolized on a culture plate, or a standard proliferation assay for T-helper cells.

Recombinant Yersinia can also be employed in an ex vivo regime for inducing CTLs specific for a protein. The procedure to develop such specific CTLs in vitro is known in the art, e.g., as disclosed by the United States Patent No. 5,342,774. Briefly, a blood sample containing T cell precursors is taken from a mammal. PBLs are purified from such blood sample and are incubated with stimulator cells expressing an antigenic epitope in the context of an MHC molecule. CTLs specific for such epitope produced can be detected by assays such as an assay for Cr⁵¹ release or secretion of IFN-gamma.

According to the present invention, a mixture of a recombinant Yersinia and an antigen presenting cell can be used as the "stimulator cell" in such an in vitro procedure for producing CTLs specific for the protein being delivered. The MHC determinants expressed by the antigen presenting cell

used are compatible with those expressed by the mammal from which PBLs are isolated, and at least one of these MHC molecules is capable of presenting, to T cells, one or more epitopes derived from the protein 5 being delivered. CTL cells generated as such can be administered, in a therapy regimen of adoptive transfer, to a mammal a pathological condition characterized by an abnormal expression of the protein used in the delivery system. See teachings by 10 Greenberg (1986) J. Immunol. 136 (5): 1917; Riddel et al. (1992) Science 257: 238; Lynch et al. (1991) Eur. J. Immunol. 21: 1403; and Kast et al. (1989) Cell 59: 603 for adoptive transfer. CTLs, by lysing the cells abnormally expressing such antigens, can alleviate or 15 treat the pathological condition at issue such as a tumor, an infection with a parasite or a virus.

Accordingly, the present invention contemplates methods and compositions for treating pathological conditions. The pathological conditions contemplated by the present invention include tumors and infections by pathogens such as bacteria, parasites, fungus or virus.

20

25

30

By "treating", is meant alleviating or inhibiting a pathological condition, e.g., inhibiting tumor growth or metastasis, reducing the size of tumor, or diminishing symptoms of a pathogen infection.

The recombinant Yersinia of the present invention can also be employed in vivo, i.e., introducing recombinant Yersinia into a mammal, such as a human or rodent subject.

For in vivo use of recombinant Yersinia, the safety can be tested in animals beforehand. case, the recombinant Yersinia may be administered to the animal orally or directly into the stomach. animals may be sacrificed a few days (1-3 days) after the administration of the recombinant Yersinia. intestines are washed and the Peyer patches or the faeces can be examined for viable Yersinia. e.g., Sory et al. (1992) Infect. Immun. 60: 3830-3836. The recombinant Yersinia may also be administered to the animal by intraperitoneal injection. Organs of sacrificed animals such as spleen and liver can be examined for the presence of intracellular Yersinia, an indication of insufficient safety. Intracellular Yersinia may be detected by e.g., cultivating cell extracts on solid medium. See teachings by Sory et al. (1988) Microb. Pathogen 4: 431-442.

5

10

15

20

25

30

A safe recombinant Yersinia may be employed in an immunogenic composition to induce an immune response for treating various pathological conditions in mammals. The pathological conditions contemplated by the present invention include tumors and pathogen infections, as disclosed herein.

The immunogenic compositions can include, in addition to a recombinant Yersinia, other substances such as cytokines, adjuvants and pharmaceutically acceptable carriers. Cytokines can also be included in such immunogenic composiitons using additional recombinant Yersinia of the present invention capable of delivering a cytokine, for example.

These immunogenic compositions may be administered to the subject in any convenient manner, such as orally, intraperitoneally, intravenously or subcutaneously. Specific immune responses induced by such compositions can lead to the CTL-mediated or antibody-mediated killing of the pathogens or cells with abnormal expression of a relevant antigen, thus alleviating the relevant pathological condition.

5

10

15

20

The present invention is further illustrated by the following examples.

All the publications mentioned in the present disclosure are incorporated herein by reference. The terms and expressions which have been employed in the present disclosure are used as terms of description and not of limitation, and there is no intention in the use of such terms and expressions of excluding any equivalents of the features shown and described or portions thereof, it being recognized that various modifications are possible within the scope of the invention.

Example 1

Bacterial Strains, Plasmids and Growth Conditions

The work was carried out with

Y.enterocolitica E40 (pYV40) (see, M.P. Sory et al.
(1995) "Identification of the YopE and YopH domains required for secretion and internalization into the cytosol of macrophages, using the cyaA gene fusion approach" Proc. Nat'l Acad. Sci. USA 92: 11998-12002), its isogeneic ampicillin sensitive derivative

MRS40(pYV40) (see, M.R. Sarker et al., and their various non-polar mutants. Plasmids are listed in Table 1. Bacteria were grown in Brain Heart Infusion (BHI) (Difco, Destroit, Michigan). After overnight

preculture, bacteria were diluted 1/20 in fresh BHI, allowed to grow for 30 minutes at room temperature, and synthesis of the Yop virulon was induced by incubation for 150 minutes at 37°C before infection.

Table 1: Exemplary Antigens

	Gene	мнс	Peptide	Position	SEQ ID
	MAGE-1	HLA-A1	EADPTGHSY	161-169	1
5		HLA-Cw16	SAYGEPRKL	230-238	2
	MAGE - 3	HLA-A1	EVDPIGHLY	168-176	3
		HLA-A2	FLWGPRALV	271-279	4
		HLA-B44	MEVDPIGHLY	167-176	5
	BAGE	HLA-Cw16	AARAVFLAL	2-10	6
10	GAGE-1,2	HLA-Cw16	YRPRPRRY	9-16	7
	RAGE	HLA-B7	SPSSNRIRNT	11-20	8
	GnT-V	HLA-A2	VLPDVFIRC(V)	2-10/11	9
	MUM - 1	HLA-B44	EEKLIVVLF	exon 2/	10
				intron	
			EEKLSVVLF		11
			(wild type)		
15	CDK4	HLA-A2	ACDPHSGHFV	23-32	12
			ARDPHSGHFV		13
			(wild type)]
	β-catenin	HLA-A24	SYLDSGIHF	29-37	14
			SYLDSGIHS		15
			(wild type)		
	Tyrosinase	HLA-A2	MLLAVLYCL	1-9	16
20		HLA-A2	YMNGTMSQV	369-377	17
		HLA-A2	YMDGTMSQV	369-377	18
		HLA-A24	AFLPWHRLF	206-214	19

	HLA-B44	SEIWRDIDF	192-200	20
	HLA-B44	YEIWRDIDF	192-200	21
	HLA-DR4	QNILLSNAPLGPQ FP	56-70	22
	HLA-DR4	DYSYLQDSDPDSF QD	448-462	23
Melan-A ^{MART-1}	HLA-A2	(E) AAGIGILTV	26/27-35	24
	HLA-A2	ILTVILGVL	32-40	25
gp100 ^{Pmel117}	HLA-A2	KTWGQYWQV	154-162	26
	HLA-A2	ITDQVPFSV	209-217	27
	HLA-A2	YLEPGPVTA	280-288	28
	HLA-A2	LLDGTATLRL	457-466	29
	HLA-A2	VLYRYGSFSV	476-485	30
DAGE	HLA-A24	LYVDSLFFL	301-309	31
MAGE-6	HLA-Cw16	KISGGPRISYPL	292-303	32

5

10

Example 2 Construction of the Polymutant Strains

To construct the yopHOPEM polymutant strain, the yopE, yopH, yopO, yopM and yopP genes were 5 successively knocked out by allelic exchange in the MRS40 strain using the suicide vectors pMRS101 and pKNG101. See, K. Kaniga et al. (1991) "A wide-host range suicide vector for improving reverse genetics in gram-negative bacteria: inactivation of the blaA gene 10 of Yersinia enterocolitica" Gene 109: 137-141 and M.R. Sarker et al. (1997) "An improved version of suicide vector pKNG101 for gene replacement in Gramnegative bacteria" Mol. Microbiol. 23: 409-411. various deletions are described in Table 2 in the 15 "suicide vectors and mutators" section. The YopE gene was first mutated using the mutator pPW52 (see, P. Wattiau et al. (1993) "SycE, a chaperone-like protein of Yersinia enterocolitica involved in the secretion of YopE" Mol. Microbiol. 8: 123-131), giving strain 20 MRS40(pAB4052). Mutation of the YopH gene in this strain with the mutator pAB31 (see, S.D. Mills et al. (1997) "Yersinia enterocolitica induces apoptosis in macrophages by a process requiring functional type III secretion and translocation mechanisms and involving 25 YopP, presumably acting as an effector protein" Proc. Natl. Acad. Sci. USA 94: 12638-12643) gave the double yopEH mutant MRS40(pAB404). The triple yopEHO mutant MRS40(pAB405) was then obtained by allelic exchange with the mutator pAB34 (see, S.D. Mills et al., 1997). 30 The YopP gene was then mutated with mutator pMSK7 (see

S.D. Mills et al. (1997)), leading to the yopEHOP mutant MRS40(pMSK46). The yopHOPEM strain MRS40(pABL403) was finally obtained by allelic exchange with the yopM mutator pAB38 (see, S.D. Mills et al., 1997).

5

Table 2: Plasmids

Plasmids Relevant		References		
_	Characteristics			
pYV				
pABL403	pYV40 yopE ₂₁ , yopH $_{\Delta}^{1\cdot352}$ yopO $_{\Delta}^{65\cdot558}$, yopP ₂₃ , yopM ₂₃	see Example 2 of the present specification		
	Suicide Vectors and mutators			
pKNG101	oriR6K sacBR+ onTRK2 strAB+	K. Kaniga et al. (1991) Gene 109: 137-141.		
pMRS101	oriR6K sacBR+ onTRK2 strAB+ oriColE1 bla +	M.R. Sarker and G.R. Cornelis (1997) Mol. Microbiol. 23: 409-411.		
pAB31	pMRS101 yopHa ₁₋₃₅₂ +	S.D. Mills et al. (1997) Proc. Natl. Acad. Sci. USA 94: 12638-12643.		
pAB34	pMRS101 yopO ₄₆₅₋₅₅₈ +	S.D. Mills et al. (1997)		
pAB38	pMRS101 yopM ₂₃ +	S.D. Mills et al. (1997)		
pMSK7	pMRS101 yopP ₂₃ +	S.D. Mills et al. (1997)		
pPW52	pKNG101 yopE ₂₁ +	P. Waattiau and G.R. Cornelis (1993) Mol. Microbiol. 8: 123-131.		

5

Example 3

Construction of a Plasmid Encoding YopE₁₃₀-MAGE-1 and Introduction of this Plasmid into Yersinia

The sequence encoding protein MAGE-1 was inserted in frame with a sequence encoding a truncated YopE, YopE₁₃₀, containing the first 130 amino acids of YopE. Such a plasmid is graphically depicted in Figure 1.

The open reading frame of MAGE-1 was amplified by PCR using a MAGE-1 cDNA cloned in pcDNAI/Amp (Invitrogen, Carlsbad, California) as template. The upstream primer,

AAACTGCAGATGTCTCTTGAGCAGAGGAGTC, consisted of the
first nucleotides of the open reading frame of MAGE-1
preceded by a PstI site. The downstream primer,
AAACTGCAGTCAGACTCCCTCTTCCTCCTC, consisted of
nucleotides complementary to the last nucleotides of
the open reading frame of MAGE-1 followed by a PstI

site. The PCR product was digested with PstI and inserted in frame with the truncated YopE at the PstI site of vector pMS111 (see, Sory et al. (1994)

Molecular Microbiology 14: 583-594). pMS111-MAGE-1 was electroporated in bacteria strain DH5αF'IQ. DNA

was extracted from some clones and the DNA of a positive recombinant clone was electroporated in bacteria strain SM10. After mobilization of pMS111 from SM10 in Yersinia MRS40 (pABL403), recombinant clones were then selected on agar-containing medium,

supplemented with nalidixic acid, sodium-arsenite and chloramphenicol. MRS40 is an isogeneic derivative of

E40 sensitive to ampicillin (see, Sory et al. (1995) Proc. Natl. Acad. Sci. USA 92: 11998-12002).

Example 4 Targeting EBV-Transformed B Cells

One colony of Yersinia MRS40 (pABL403) 5 containing pMS111-MAGE-1 was then grown overnight at 28°C in LB medium supplemented with nalidixic acid, sodium m-arsenite and chloramphenicol. The overnight culture was diluted in fresh medium in order to obtain an OD (optical desity) of 0.2. The fresh culture was amplified at 28°C for approximately 2 hours. 10 bacteria were washed in 0.9% NaCl and resuspended at 108 bacteria per ml in 0.9% NaCl. 50,000 EBVtransformed HLA-A1 B cells (KASOII-EBV) were placed in microwells (96 wells round-bottomed) and pelleted 15 by centrifugation. The supernatant was discarded and various dilutions of bacteria were added in 100 ul of complete RPMI 1640 (culture media was supplemented with 10% FCS and with L-arginine (116 mg/ml), Lasparagine (36 mg/ml), L-glutamine (216 mg/ml). Two 20 hours after infection, gentamicin (30 $\mu g/ml$) was added for the next two hours, and the cells were finally washed three times.

As a negative control, the same cells were also infected with Yersinia~MRS40~(pABL403) containing pMS621, a plasmid which encodes only the truncated YopE, i.e., $YopE_{130}$.

25

Example 5 Recognition of Targeted B Cells by MZ2-CTL 82/30

MZ2-CTL 82/30 are specific for the MAGE-1 5 peptide EADPTGHSY which is presented by HLA-A1 (U.S. Patent No. 5,342,774). 5000 MZ2-CTL 82/30 cells were added in each microwell containing the Yersinia in a final volume of 100 μ l of Iscove's complete medium (culture medium was supplemented with 10% human serum, 10 L-arginine (116 mg/ml), L-asparagine (36 mg/ml), Lglutamine (216 mg/ml), streptomycine (0.1 mg/ml), penicillin (200 U/ml), IL-2 (25 U/ml) and gentamicin (15 μ g/ml). After overnight incubation, the presence of IFN-gamma (that is produced by CTL upon activation) 15 in the supernatant of the co-culture was tested in a standard ELISA assay (Biosource, Fleurus, Belgium). Figure 2A graphically depicts such a procedure. As indicated in Figure 2B, the HLA-A1 B cells infected with Yersinia encoding YopE, 30 - MAGE-1 20 were recognized by the CTL 82/30, while the same cells infected with the control plasmid $YopE_{130}$ were not. The optimal concentration of bacteria is around

1,000,000 per microwell.

WHAT IS CLAIMED IS:

1. A mutant Yersinia strain comprising at
least one mutation in at least one Yersinia effector-
encoding gene, whereby said mutant Yersinia strain is
deficient in the production of at least one functional
effector protein.

- 2. The mutant Yersinia strain according to claim 1, wherein said effector encoding gene is selected from the group consisting of YopE, YopH, YopO, YopM and YopP of Y. enterocolitica; and YopE, YopH, YpkA, YopM and YopJ of Y. pseudotuberculosis.
 - 3. The mutant Yersinia strain according to claim 1, wherein said mutation is a mutation of the promoter sequence of said effector gene.
- 4. The mutant Yersinia strain according to claim 1, wherein said mutation is a mutation of the coding sequence of said effector gene.
- 5. A quintuple mutant Yersinia strain having the designation of Y. enterocolitica yopEHOMP or Y. pseudotuberculosis yopEHAOJ.
- 6. The quintuple mutant Yersinia strain according to claim 5, having the designation of Yersinia enterocolitica MRS40(pABL403).

1	 An expression vector for delivering a
2	heterologous protein into a eukaryotic cell, which
3	comprises in the 5' to 3' direction:
4	a promoter from a Yersinia virulon gene;
- 5	a first DNA sequence encoding a delivery
6	signal from a Yersinia effector protein, operably
7	linked to said promoter; and
8	a second DNA sequence coding for said
9	heterologous protein, fused in frame to the 3' end of
10	said first DNA sequence.
1	8. The expression vector of claim 7,
2	wherein said Yersinia virulon gene is a Yersinia
3	effector-encoding gene.
1	9. The expression vector of claim 8,
2	wherein said effector-encoding gene is selected from
3	the group consisting of YopE, YopH, YopO, YopM and
4	YopP of Y. enterocolitica; and YopE, YopH, YpkA, YopM
5	and YopJ of Y. pseudotuberculosis.
1	10. The expression vector of claim 9,
2	wherein said effector-encoding gene is Y.
3	enterocolitica of YopE.
1	11. The expression vector of claim 7,
2	wherein said effector protein is selected from the
3	group consisting of YopE, YopH, YopO, YopM and YopP of
4	Yersinia enterocolitica; and YopE, YopH, YpkA, YopM
5	and YopJ of Y. pseudotuberculosis.

_____WO 99/45098 PCT/IB99/00587

1	12. The expression vector of claim 11,
2	wherein said effector protein is one of Yersinia
3	enterocolitica YopE or Y. pseudotuberculosis YopE.
1	13. The expression vector of claim 7,
2	wherein said delivery signal is Y. enterocolitica
3	YopE ₁₃₀ .
1	14. The expression vector of claim 7,
2	wherein said heterologous protein comprises at least
3	one epitope of a naturally occurring protein.
1	15. The expression vector of claim 14,
2	wherein said naturally occurring protein is a tumor
3	associated protein or a pathogen antigen.
1	16. The expression vector of claim 15,
2	wherein said tumor associated protein is selected from
3	the group consisting of members of the MAGE family,
4	the BAGE family, the DAGE/Prame family, the GAGE
5	family, the RAGE family, the SMAGE family, NAG,
6	Tyrosinase, Melan-A/MART-1, gp100, MUC-1, TAG-72,
7	CA125, p21ras, p53, HPV16 E7, HOM-MEL-40, HOM-MEL-55,
8	NY-COL-2, HOM-HD-397, HOM-RCC-1.14, HOM-HD-21, HOM-
9	NSCLC-11, HOM-MEL-2.4, and HOM-TES-11.
1	17. The expression vector of claim 16,
2	wherein said tumor-associated protein is MAGE-1.
1	18. A Yersinia of a Yersinia mutant strain
2	of any one of claims 1-6 for delivery of a

3	heterologous protein into a eukaryotic cell, wherein
4	said Yersinia is transformed with an expression vector
5	which comprises in the 5' to 3' direction:
6	a promoter from a Yersinia virulon gene;
1	a first DNA sequence encoding a delivery
2	signal from a Yersinia effector protein, operably
3	linked to said promoter; and
4	a second DNA sequence coding for said
5	heterologous protein, fused in frame to the 3' end of
6	said first DNA sequence.
1	19. The Yersinia according to claim 18,
2	wherein said <i>Yersinia</i> virulon gene is a <i>Yersinia</i>
3	effector-encoding gene.
1	20. The Yersinia according to claim 19,
2	wherein said effector-encoding gene is selected from
3	the group consisting of YopE, YopH, YopO, YopM and
4	YopP of Y. enterocolitica; and YopE, YopH, YpkA, YopM
5	and YopJ of Y. pseudotuberculosis.
1	21. The Yersinia according to claim 20,
2	wherein said effector-encoding gene is Y.
3	enterocolitica YopE.
1	22. The Yersinia according to claim 18,
2	wherein said effector protein selected from the group
3	consisting of YopE, YopH, YopO, YopM and YopP of
4	Yersinia enterocolitica; and YopE, YopH, YpkA, YopM
5	and VonJ of V nseudotuberculosis

1	23. The Yersinia according to claim 22,
2	wherein said effector protein is one of Yersinia
3	enterocolitica's YopE or Y. pseudotuberculosis's YopE.
1	24. The Yersinia according to claim 18,
2	wherein said delivery signal is Y. enterocolitica
3	YopE ₁₃₀ .
1	25. The Yersinia according to claim 18,
2	wherein said heterologous protein comprises at least
3	one epitope of a naturally occurring protein.
1	26. The Yersinia according to claim 25,
2	wherein said naturally occurring protein is a tumor-
3	associated protein or a pathogen antigen.
1	27. The Yersinia according to claim 26,
2	wherein said tumor-associated protein is selected from
3	the group consisting of members of the MAGE family,
4	the BAGE family, the DAGE/Prame family, the GAGE
5	family, the RAGE family, the SMAGE family, NAG,
6	Tyrosinase, Melan-A/MART-1, gp100, MUC-1, TAG-72,
7	CA125, p21ras, p53, HPV16 E7, HOM-MEL-40, HOM-MEL-55,
8	NY-COL-2, HOM-HD-397, HOM-RCC-1.14, HOM-HD-21, HOM-
9	NSCLC-11, HOM-MEL-2.4, and HOM-TES-11.
1	28. The Yersinia according to claim 27,
2	wherein said tumor-associated protein is MAGE-1.
1	29. A method for delivering a heterologous
2	protein into a eukaryotic cell, comprising contacting

3 said eukaryotic cell with a Yersinia of a mutant 4 strain of claim 1 transformed with an expression vector characterized by in 5' to 3' direction: 5 a promoter from a Yersinia virulon gene; 6 1 a first DNA sequence encoding a delivery 2 signal from a Yersinia effector protein, operably 3 linked to said promoter; and 4 a second DNA sequence coding for said 5 heterologous protein, fused in frame to the 3' end of 6 said first DNA sequence. 1 The method of claim 29, wherein said 2 eukaryotic cell is an antigen presenting cell. 1 The method of claim 30, wherein said 2 antigen presenting cell is selected from the group 3 consisting of a B cell, a macrophage, a dendritic 4 cell, a monocyte, a follicular cell, and a fibroblast. 1 32. The method of claim 30, wherein said 2 antigen presenting cell expresses an MHC molecule 3 capable of presenting one or more epitopes derived

from said heterologous protein.

4

1

2

3

4

5

l	34. The method of claim 33, wherein said
2	effector protein is one of Yersinia enterocolitica
3	yopE or Y. pseudotuberculosis YopE.
1	35. The method of claim 29, wherein said
2	effector protein is $YopE_{130}$ of Yersinia enterocolitica.
1	36. The method of claim 29, wherein said
2	heterologous protein comprises at least one epitope of
3	a naturally occurring protein.
1	37. A Yersinia according to claim 36,
2	wherein said naturally occurring protein is a tumor
3	associated protein or a pathogen antigen.
1	38. The expression vector of claim 37,
2	wherein said tumor associated protein is selected from
3	the group consisting of members of the MAGE family,
4	the BAGE family, the DAGE/Prame family, the GAGE
5	family, the RAGE family, the SMAGE family, NAG,
6	Tyrosinase, Melan-A/MART-1, gp100, MUC-1, TAG-72,
7	CA125, p21ras, p53, HPV16 E7, HOM-MEL-40, HOM-MEL-55,
8	NY-COL-2, HOM-HD-397, HOM-RCC-1.14, HOM-HD-21, HOM-
9	NSCLC-11, HOM-MEL-2.4, and HOM-TES-11.
1	39. The method of claim 38, wherein said
2	tumor-associated protein is MAGE-1.
1	40. The method of claim 29, wherein said
2	Yersinia virulon gene is a Yersinia effector-encoding
3	gene.

1	41. The method of claim 40, wherein said
2	effector-encoding gene is selected from the group
3	consisting of YopE, YopH, YopO, YopM and YopP of Y.
4	enterocolitica; and YopE, YopH, YpkA, YopM and YopJ of
5	Y. pseudotuberculosis.
1	42. The method of claim 41, wherein said
2	effector-encoding gene is YopE of Y. enterocolitica.
1	43. A method for inducing an immune
2	response specific for a heterologous protein,
3	comprising.the steps of:
4	(a) selecting an antigen presenting cell
5	expressing an MHC molecule capable of presenting at
6	least one epitope of said heterologous protein;
7	(b) forming a cell mixture by contacting
8	said antigen presenting cell with a Yersina of a
9	mutant Yersinia strain of claim 1 transformed with an
10	expression vector thereby delivering said heterologous
11	protein into said antigen presenting cell, wherein
12	said expression vector is characterized by in 5' to 3'
13	direction:
14	a promoter from a Yersinia virulon gene;
15	a first DNA sequence encoding a delivery
16	signal from a Yersinia effector protein, operably
17	linked to said promoter;
18	a second DNA sequence coding for said
19	heterologous protein, fused in frame to the 3' end of
20	said first DNA sequence; and
21	(c) contacting a sample containing
22	peripheral blood lymphocytes taken from a subject.

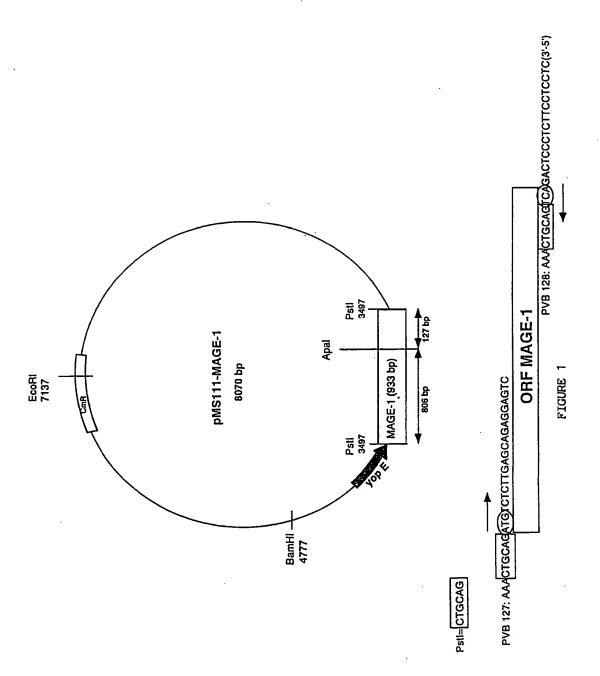
23	with the cell mixture formed in step (b) thereby
24	inducing an immune response specific for said
25	heterologous protein.
1	44. The method of claim 43, wherein said
2	epitope of is from a tumor associated protein.
1	45. The method of Claim 44, wherein said
2	tumor associated protein is selected from the group
3	consisting of members of the MAGE family, the BAGE
4	family, the DAGE/Prame family, the GAGE family, the
5	RAGE family, the SMAGE family, NAG, Tyrosinase, Melan
6	A/MART-1, gp100, MUC-1, TAG-72, CA125, p21ras, p53,
7	HPV16 E7, HOM-MEL-40, HOM-MEL-55, NY-COL-2, HOM-HD-397
8	HOM-RCC-1.14, HOM-HD-21, HOM-NSCLC-11, HOM-MEL-2.4,
9	and HOM-TES-11.
1	46. The method of claim 45, wherein said
2	tumor associated protein is MAGE-1.
1	47. The method of claim 46, wherein said
2	epitope is from MAGE-1 and said MHC molecule is HLA-
3	A1.
1	48. An immunogenic composition, comprising
2	a recombinant Yersinia according to claim 18.
1	49. A method of inducing a CTL response
2	specific for a heterologous protein in a subject in
3	need of such response, comprising the steps of:
_	

1	(a) obtaining from said subject an antigen
2	presenting cell expressing an MHC molecule;
3	(b) forming a cell mixture by contacting
4	said antigen presenting cell with a Yersina of a
5	mutant Yersinia strain of claim 1 transformed with an
6	expression vector, wherein said expression vector is
7	characterized by in 5' to 3' direction:
8	a promoter from a Yersinia virulon gene;
9	a first DNA sequence encoding a delivery
10	signal from a yersinia effector protein, operably
11	linked to said promoter; and
12	a second DNA sequence fused in frame to the
13	3' end of said first DNA sequence, wherein said second
14	DNA sequence codes for at least one epitope of said
15	heterologous protein which is presented by said MHC
16	molecule of said antigen presenting cell;
17	(c) contacting a sample containing
18	peripheral blood lymphocytes taken from said subject,
19	with the cell mixture formed in step (b), thereby
20	producing CTLs specific for said heterologous protein;
21	and
22	(d) administering CTLs produced in step (c)
23	to said subject thereby inducing a CTL response
24	specific for said heterologous protein in said
25	subject.
1	50. A method for determining the efficacy
2	of a vacciantion regimen in a subject, wherein said
3	subject is vaccinated with an antigen, comprising the
4	steps of:

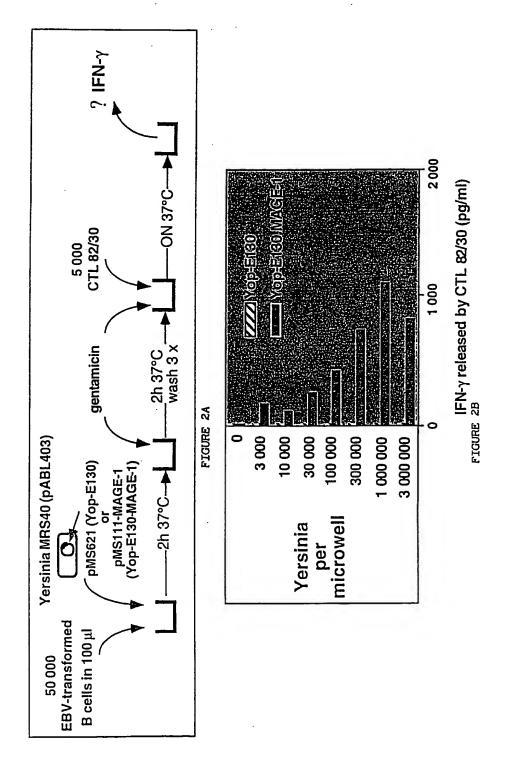
obtaining from said subject an antigen 1 presenting cell expressing an MHC molecule; 2 forming a cell mixture by contacting 3 said antigen presenting cell with a Yersina of a 4 mutant Yersinia strain of claim 1 transformed with an 5 expression vector, wherein said expression vector is 6 7 characterized by in 5' to 3' direction: 8 a promoter from a Yersinia virulon gene; a first DNA sequence encoding a delivery 9 10 signal from a yersinia effector protein, operably 11 linked to said promoter; and 12 a second DNA sequence fused in frame to the 3' end of said first DNA sequence, wherein said second 13 14 DNA sequence codes for at least one epitope of said antigen which is presented by said MHC molecule of 15 said antigen presenting cell; 16 17 (c) contacting a sample containing 18 peripheral blood lymphocytes taken from said subject, with the cell mixture formed in step (b), and assaying 19 the presence of an immune response specific for said 20 antigen thereby determining the efficacy of said 21 vaccination regimen. 22 51. A method for treating a pathological 1 2 disorder in a subject, comprising administering the versinia of claim 18 to said subject, wherein the 3 4 heterologous protein elicits an immune response

specific to said pathological disorder.

5



Yersinia which produces a YopE130.MAGE-1 fusion protein Anti-MAGE-1.A1 CTL recognize HLA-A1 cells incubated with



3/7

1 61 121 181 241 301 361 421 481 541 601	attttcgacg aaaaaaccga atgtatctaa cagttgaggc aacgaaatgc attgcctgga cggaattacc cggaattacc	aaaaactata taataaagat tacttttttg agaaaatgtt ccctccgggg ccgacaagcc tccgcattta gcagagcctg tcctttactg	tatatatata attttcagaa caagaaccat aaatctaagg aatggtgaac catgagctag gagagtttag aaatcacttc gaatatttag	tatttaatat aggcattcaa tacgtcattc ctgaatatta agaggggaat aactaaataa tggcgtcatg aagttgataa gtgccgctaa	tggataagta gtatggtttc tatgtttata ttctgattta taatgcatgg ggcggtttca tctggggctg taattctctt taacaatctg taatcagctg tgataacaat	atttgcaatg aacccaagaa actgagatgc tcggaatggg aggttacgcg agttctttgc acagaattac aaggcattat gaagaattac
661 721 781 841 901 961 1021 1081 1141 1201 1261 1321	cattacctga aattgtcaga tgaaaacatt taactgatct tttctggatt taagatcttt tgatcgaact ttgctgaagt tgagagagtt taattgatcc	tttacctcct gttgcaaaac acccgattta gccagaatta atcggaattta gccagcgtta acctgaattg tcccgatata atatgaattt	tcactggaat ttgccettct ccccettccc ccgcagagtt ccaccaaact ccccettcac cctccacgct ccgcaaaacc cctgagtcag gctcatgaga	ttettgetge tgaetgegat tgaaaacact taacettett tgtataatet tggtagaact tagaaegttt tgaaaetget tggaagatet ctatagaeaa	tgataacaat tggtaataat ttatgctgat taatgtcaga agatgttct caatgcatcc tgatgtcaga aatcgcttca acacgtagag tcggatggac acttgaagat aatatcatta	cagctggaag aacaattcac gaaaattatt gataatattt agcaatgaaa gataatcagt tttaatcatc tacaacgctc tctgaacgtg gatgtatttg

4/7

_						
1	gaattcccca	actttgacac	cgataaccgg	ttcaatagta	tctggaatag	acagcgaaag
61	ttgttgaaat	aattgagtga	tagcttgttc	aaatgaatac	atttgatctc	ctaatagtta
121	gataaaatat	caacttaacc	aaagcactct	cggcagacca	tcaattttag	cctataattt
181	ttagttttta	ttttgtctaa	tataacaaca	aaaacagcag	cggtttttta	tataaccacc
241	ggctattttc	ccactaagat	aaccttgttt	taatagccaa	gggaataaat	agtcatgaaa
301	atatcatcat	ttatttctac	atcactgccc	ctgccggcat	cagtgtcagg	atctagcagc
361	gtaggagaaa	tgtctgggcg	ctcagtctca	cagcaaaaaa	gtgatcaata	tgcaaacaat
421	ctggccgggc	gcactgaaag	ccctcagggt	tccagcttag	ccagccgtat	cattgagagg
481	ttatcatcaa	tggcccactc	tgtgattgga	tttatccaac	gcatgttctc	ggaggggagc
541	cataaaccgg	tggtgacacc	agcactcacg	cctgcacaaa	tgccaagccc	tacgtctttc
601	agtgatagta	tcaagcaact	tgctgctgag	acgctgccaa	aatacatgca	gcagttgagt
661	agcttggatg	cagagacgct	gcagaaaaat	catgaccagt	tcgccacggg	cagcggccct
721	cttcgtggca	gtatcactca	atgccaaggg	ctgatgcagt	tttgtggtgg	ggaattgcaa
781	gctgaggcca	gtgccatttt	aaacacgcct	gtttgtggta	ttcccttctc	gcagtgggga
841	actgttggtg	gggcggccag	cgcgtacgtc	gccagtggcg	ttgatctaac	gcaggcagca
901	aatgagatca	aagggctggg	gcaacagatg	cagcaattac	tgtcattgat	gtgatatgga
961	taaaaacaag	ggggtagtgt	ttcccccttt	ttctatcaat	attgcgaata	tcttcgtccc
1021	tgatctttca	ggggcgaatc	gttttttagc	atgctcattg	ttagaatttc	tgacttatct
1081	ctcttctgta	ttactactca	tactctggaa	aatcctgagc	atttatatct	atggattgat
1141						

1	agggcattgg	aattaaaaat	atatttatct	aaatgatgat	gagtttaaat	tacatttgcg
61	tattaaaatg	aataacgcat	tattaacgta	ttaccatctg	ttcccgctta	attttttaaa
121	aaatttaagg	taacaatgag	tatatatctt	atgggaaaag	ccaaaaaact	aacgaacact
181	ataataattc	gattaacatc	aatgaaaata	cacggctcac	ctattattaa	aataatacga
241	ctagcattat	aagaaaaaat	attttttatg	tttatagtat	aggcgtgtat	ttaattagtt
301	cttaatttaa	ttaaggaggg	aagcatgaac	ttatcattaa	gcgatcttca	tcqtcaqqta
361	tctcgattgg	tgcagcaaga	gagcggtgat	tgtaccggga	aattaagagg	taacqttqct
421	gccaataaag	aaactacctt	tcaaggtttg	accatagcca	gtggtgccag	agagtcagaa
481	aaagtatttg	ctcaaactgt	actaagccac	gtagcaaata	ttgttctaac	tcaagaagat
541	accgctaagc	tattgcaaag	cacggtaaag	cataatttga	ataattatga	attaagaagt
601	gtcggcaatg	gtaatagtgt	acttgtcagt	ttacgtagtg	accaaatgac	actacaagac
661	gccaaagtgc	tgttggaggc	tgcattgcga	caagagtcgg	gagcgagggg	gcatgtatca
721	tctcattcac	attcagtcct	tcacgcaccg	ggaaccccgg	tgcgtgaagg	actgcgttca
781				ccgcgtgaac		
841	cacggggctg	gcgaagccag	agccaccgca	ccaagcactg	tttctcctta	tggcccagaa'
901				acattgcgca		
961		gttacttaca	agcctgcggc	ggtgaaaagc	taaaccgatt	tagagatatt
1021				gatcttaatg		
1081		ccatagcgtg	ccagtatccg	ctacaatctc	aacttgaaag	ccatttccgt
1141	atgctggcag	aaaaccgaac	gccagtgttg	gctgttttag	cgtccagttc	tgagatagcc
1201	aatcaaagat	tcggtatgcc	agattatttc	cgccagagtg	gtacctatgg	cagtatcact
1261	gtagagtcta	aaatgactca	gcaagttggt	ctcggtgacg	ggattatggc	agatatgtat
	actttaacga					
	aattggcccg					
	gatcaaacag					
	gatgactcca					
	ctgattggcg					
	gtcagccaaa					
	gttctgatta					
	ttattcctat					
	ctttaaacct					
1861	aacaaaaata	taaacaacaa	aattaaaagt	tatgtgtcta	cttttacttt	atgtaaccaa
1921	acccattaat	ggataccgta	cgtttttctt	ttatagaatt	aaaccagtaa	atgagatgat
1981	gaaggacgat	gatcatcgtc				

...WO 99/45098

6/7

					+-+	agagagagat
1	atgattgggc	caatatcaca	aataaacagc	tteggtgget	tatcagaaaa	agagacccgc
61	totttaatca	gtaatgaaga	gcttaaaaat	atcataatac	agttggaaac	tgatatagcg
121	gatggatect	ggttccataa	aaattattca	cacctagata	tagaagtcat	gcccgcatta
	atosattosaa	cgaacaataa	atateceraa	atreatrtta	attttqttac	atctccccag
181	gtaattcagg	cgaacaacaa	acacccggaa	acguacece		t-
241	gacctttcga	tagaaataaa	aaatgtcata	gaaaatggag	ttggatette	Cogollcala
3.01	attaacatgg	gggagggtgg	aatacatttc	agtgtaattg	attacaaaca	tataaatggg
361	aaaacatctc	tgatattatt	tgaaccagta	aactttaata	gtatggggcc	agcgatactg
301	aaaacacccc				atmattacca.	tttttccato
421	gcaataagta	caaaaacggc	cattgaacgt	tateaattae	Clyattycta	cccccaog
481	gtggaaatgg	atattcagcg	aagctcatct	gaatgtggta	tttttagttt	ggcactggca
541	aaaaaacttt	acaccgagag	agatageetg	ttgaaaatac	atgaagataa	tataaaaggt
601	atattaagtg	atagtgaaaa	teetttacce	cacaataagt	togatccgta	tctcccggta
907	atattaagtg	acagegaaaa				tagtaagggg
661	actttttaca	aacatactca	aggtaaaaaa	cgtcttaatg	aatatttaaa	Lactaatteg
721	cagggagttg	gtactgttgt	taacaaaaaa	aatgaaacca	tctttaatag	gtttgataac
781	aataaatcca	ttatagatgg	aaaggaatta	tcagtttcgg	tacataaaaa	gagaatagct
		cacttctcaa				

7/7

 ${\tt CAGTGGTTTTCTCTCTTTCGAGAAGGGGTTGGTAAGATCTTTTCGGGGGAAGATGTTTAACTTTTCAATTGCTCGT}$ AACCTTACTGAGACACTCCATGCAGCCCAGAAAACGACTTCGCAGGAGCTAAGGTCTGATATCCCCAATGCTCTCA GTAATCTCTTTGGAGCCAAGCCACAGACCGAACTGCCGCTGGGTTGGAAAGGGAAGCCTTTGTCAGGAGCTCCGGA TCTTGAAGGGATGCGAGTGGCTGAAACCGATAAGTTTGCCGAGGGCGAAAGCCATATTAGTATAATAGAAACTAAG GATAATCAGCGGTTGGTGGCTAAGATTGAACGCTCCATTGCCGAGGGGCATTTGTTCGCAGAACTGGAGGCTTATA AACACATCTATAAAACCGCGGGCAAACATCCTAATCTTGCCAATGTCCATGGCATGGCTGTGGTGCCATACGGTAA CCGTAAGGAGGAAGCATTGCTGATGGATGAGGTGGATGGTTGGCGTTGTTCTGACACACTAAGAAGCCTCGCCGAT AGCTGGAAGCAAGGAAAGATCAATAGTGAAGCCTACTGGGGAACGATCAAGTTTATTGCCCATCGGCTATTAGATG TAACCAATCACCTTGCCAAGGCAGGGATAGTACATAACGATATCAAACCCGGTAATGTGGTATTTGACCGCGCTAG $\tt CGGAGAGCCCGTTGTCATTGATCTAGGATTACACTCTCGTTCAGGGGAACAACCTAAGGGGTTTACAGAATCCTTC$ AAAGCGCCGGAGCTTGGAGTAGGAAACCTAGGCGCATCAGAAAAGAGCGATGTTTTTCTCGTAGTTTCAACCCTTC TACATGGTATCGAAGGTTTTGAGAAAGATCCGGAGATAAAGCCTAATCAAGGACTGAGATTCATTACCTCAGAACC AGCGCACGTAATGGATGAGAATGGTTACCCAATCCATCGACCTGGTATAGCTGGAGTCGAGACAGCCTATACACGC TTCATCACAGACATCCTTGGCGTTTCCGCTGACTCAAGACCTGATTCCAACGAAGCCAGACTCCACGAGTTCTTGA GCGACGGAACTATTGACGAGGAGTCGGCCAAGCAGATCCTAAAAGATACTCTAACCGGAGAAATGAGCCCATTATC TACTGATGTAAGGCGGATAACACCCAAGAAGCTTCGGGAGCTCTCTGATTTGCTTAGGACGCATTTGAGTAGTGCA GCAACTAAGCAATTGGATATGGGGGTGGTTTTGTCGGATCTTGATACCATGTTGGTGACACTCGACAAGGCCGAAC GCGAGGGGGGAGTAGACAAGGATCAGTTGAAGAGTTTTAACAGTTTGATTCTGAAGACTTACAGCGTGATTGAAGA CTATGTCAAAGGCAGAGAAGGGGATACCAAGAGTTCCAGTGCGGAAGTATCCCCCTATCATCGCAGTAACTTTATG CTATCGATCGCCGAACCTTCACTGCAGAGGATCCAAAAGCATCTGGACCAGACACACTCTTTTTCTGATATCGGTT CACTAGTGCGCGCACATAAGCACCTGGAAACGCTTTTAGAGGTCTTAGTCACCTTGTCACCGCAAGGGCAGCCCGT GTCCTCTGAAACCTACAGCTTCCTGAATCGATTAGCTGAGGCTAAGGTCACCTTGTCGCAGCAATTGGATACTCTC CAGCAGCAGGAGAGTGCGAAACGGCAACTATCTATTCTGATTAATCGTTCAGGTTCTTGGGCCGATGTTGCTC GTCAGTCCCTGCAGCGTTTTGACAGTACCCGGCCTGTAGTGAAATTCGGCACTGAGCAGTATACCGCAATTCACCG TCAGATGATGGCGGCCCATGCAGCCATTACGCTACAGGAGGTATCGGAGTTTACTGATGATATGCGAAACTTTACA AGTTGCGAGACGTGACGACCATCGCCGAGCGACTGAACCGGTTGGAGCGGGAATGGATGTGA

SEQUENCE LISTING

<110> van der Bruggen, Pierre Cornelis, Guy R.

<120> DELIVERY OF PROTEINS INTO EUKARYOTIC CELLS
WITH RECOMBINANT YERSINIA

<130> 11154

<140> US 09/036,582

<141> 1998-03-06

<160> 39

<170> PatentIn Ver. 2.0

<210> 1

<211> 9

<212> PRT

<213> Human MAGE-1 peptide

<400> 1

Glu Ala Asp Pro Thr Gly His Ser Tyr 5

<210> 2

<211> 9

<212> PRT

<213> Human MAGE-1 peptide

<400> 2

Ser Ala Tyr Gly Glu Pro Arg Lys Leu 5

<210> 3

<211> 9

<212> PRT

<213> Human MAGE-3 peptide

<400> 3

Glu Val Asp Pro Ile Gly His Leu Tyr 5

<210> 4

<211> 9

<212> PRT

<213> Human MAGE-3 peptide

<400> 4

Phe Leu Trp Gly Pro Arg Ala Leu Val

<210> 5

<211> 10

<212> PRT

<213> Human MAGE-3 peptide

<400> 5

<210> 6

<211> 9

<212> PRT

<213> Human BAGE peptide

<400> 6

Ala Ala Arg Ala Val Phe Leu Ala Leu 5

<210> 7

<211> 8

<212> PRT

<213> Human GAGE-1,2 peptide

<400> 7

Tyr Arg Pro Arg Pro Arg Arg Tyr 5

<210> 8

<211> 10

<212> PRT

<213> Human RAGE peptide

<400> 8

Ser Pro Ser Ser Asn Arg Ile Arg Asn Thr 5 10

<210> 9

<211> 10

<212> PRT

<213> Human GnT-V peptide

<400> 9

Val Leu Pro Asp Val Phe Ile Arg Cys Val 5 10

<210> 10

<211> 9

<212> PRT

<213> Human MUM-1 peptide

<400> 10

Glu Glu Lys Leu Ile Val Val Leu Phe 5

<210> 11

<211> 9

<212> PRT

<213> Human MUM-1 peptide

<400> 11

Glu Glu Lys Leu Ser Val Val Leu Phe 5

<210> 12

<211> 10

<212> PRT

<213> Human CDK4 peptide

<400> 12

Ala Cys Asp Pro His Ser Gly His Phe Val 5

<210> 13

<211> 10

<212> PRT

<213> Human CDK4 peptide

<400> 13

Ala Arg Asp Pro His Ser Gly His Phe Val 5 10

<210> 14

<211> 9

<212> PRT

<213> Human β -catenin peptide

<400> 14

Ser Tyr Leu Asp Ser Gly Ile His Phe 5

<210> 15

<211> 9

<212> PRT

<213> Human β -catenin peptide

<400> 15

Ser Tyr Leu Asp Ser Gly Ile His Ser 5

<210> 16

<211> 9

<212> PRT

<213> Human Tyrosinase peptide

<400> 16

Met Leu Leu Ala Val Leu Tyr Cys Leu 5

<210> 17

<211> 9

<212> PRT

<213> Human Tyrosinase peptide

<400> 17

Tyr Met Asn Gly Thr Met Ser Gln Val 5

<210> 18

<211> 9

<212> PRT

<213> Human Tyrosinase peptide

<400> 18

Tyr Met Asp Gly Thr Met Ser Gln Val 5

<210> 19

<211> 9

<212> PRT

```
<213> Human Tyrosinase peptide
```

<400> 19

Ala Phe Leu Pro Trp His Arg Leu Phe 5

<210> 20

<211> 9

<212> PRT

<213> Human Tyrosinase peptide

<400> 20

Ser Glu Ile Trp Arg Asp Ile Asp Phe 5

<210> 21

<211> 9

<212> PRT

<213> Human Tyrosinase peptide

<400> 21

Tyr Glu Ile Trp Arg Asp Ile Asp Phe 5

<210> 22

<211> 15

<212> PRT

<213> Human Tyrosinase peptide

<400> 22

Gln Asn Ile Leu Leu Ser Asn Ala Pro Leu Gly Pro Gln Phe Pro 5 10 15

<210> 23

<211> 15

<212> PRT

<213> Human Tyrosinase peptide

<400> 23

Asp Tyr Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser Phe Gln Asp 5 10

<210> 24

<211> 10

<212> PRT

<213> Human Melan-A^{MART-1} peptide

<400> 24

Glu Ala Ala Gly Ile Gly Ile Leu Thr Val 5 10

<210> 25

<211> 9

<212> PRT

<213> Human Melan- A^{MART-1} peptide

<400> 25

Ile Leu Thr Val Ile Leu Gly Val Leu 5

<210> 26

<211> 9

<212> PRT

<213> Human gp100^{Pmel117} peptide

<400> 26

Lys Thr Trp Gly Gln Tyr Trp Gln Val

<210> 27

<211> 9

<212> PRT

<213> Human gp $100^{\text{Pme}1117}$ peptide

<400> 27

Ile Thr Asp Gln Val Pro Phe Ser Val

<210> 28

<211> 9

<212> PRT

<213> Human gp100^{Pme1117} peptide

<400> 28

Tyr Leu Glu Pro Gly Pro Val Thr Ala 5

<210> 29

<211> 10

<212> PRT

<213> Human gp100 Pmel117 peptide

<400> 29

Leu Leu Asp Gly Thr Ala Thr Leu Arg Leu 5 10

<210> 30

<211> 10

<212> PRT

<213> Human gp100^{Pmel117} peptide

<400> 30

Val Leu Tyr Arg Tyr Gly Ser Phe Ser Val 5 . 10

<210> 31

<211> 9

<212> PRT

<213> Human DAGE peptide

<400> 31

Leu Tyr Val Asp Ser Leu Phe Phe Leu 5

<210> 32

<211> 12

<212> PRT

<213> Human MAGE-6 peptide

<400> 32

Lys Ile Ser Gly Gly Pro Arg Ile Ser Tyr Pro Leu 5 10

<210> 33

<211> 1330

<212> DNA

<213> Yersinia enterocolitica

<400> 33

AAAAATGGCC AAAAACTTTC AATGGTAGAA GAGCTAAATT TGGATAAGTA ACGCATAAAA 60
ATTTTCGACG AAAAACTATA TATATATATA TATTTAATAT GTATGGTTTC ATTTGCAATG 120
AAAAAAACCGA TAATAAAGAT ATTTCAGAA AGGCATTCAA TATGTTTATA AACCCAAGAA 180
ATGTATCTAA TACTTTTTG CAAGAACCAT TACGTCATTC TTCTGATTTA ACTGAGATGC 240
CAGTTGAGGC AGAAAATGTT AAATCTAAGG CTGAATATTA TAATGCATGG TCGGAATGGG 300
AACGAAATGC CCCTCCGGGG AATGGTGAAC AGGGGGAAT GGCGGTTTCA AGGTTACGCG 360
ATTGCCTGGA CCGACAAGCC CATGAGCTAG AACTAAATAA TCTGGGGCTG AGTTCTTTGC 420
CGGAATTACC TCCGCATTTA GAGAGTTTAG TGGCGTCATG TAATTCTCTT ACAGAATTAC 480
CGGAATTGCC GCAGAGCCTG AAATCACTTC AAGTTGATAA TAACAATCTG AAGGCATTAT 540
CCGATTTACC TCCTTTACTG GAATATTTAG GTGCCGCTAA TAATCAGCTG GAAGAATTAC 600
CAGAGTTGCA AAACTCGTCC TTCTTGACAT CTATTGATGT TGATAACAAT TCACTGAAAA 660

CATTACCTGA	TTTACCTCCT	TCACTGGAAT	TTCTTGCTGC	TGGTAATAAT	CAGCTGGAAG	720
AATTGTCAGA	GTTGCAAAAC	TTGCCCTTCT	TGACTGCGAT	TTATGCTGAT	AACAATTCAC	780
TGAAAACATT	ACCCGATTTA	CCCCCTTCCC	TGAAAACACT	TAATGTCAGA	GAAAATTATT	840
TAACTGATCT	GCCAGAATTA	CCGCAGAGTT	TAACCTTCTT	AGATGTTTCT	GATAATATTT	900
TTTCTGGATT	ATCGGAATTG	CCACCAAACT	TGTATAATCT	CAATGCATCC	AGCAATGAAA	960
${\tt TAAGATCTTT}$	ATGCGATTTA	CCCCCTTCAC	TGGTAGAACT	TGATGTCAGA	GATAATCAGT	1020
TGATCGAACT	GCCAGCGTTA	CCTCCACGCT	TAGAACGTTT	AATCGCTTCA	TTTAATCATC	1080
${\tt TTGCTGAAGT}$	${\tt ACCTGAATTG}$	CCGCAAAACC	TGAAACTGCT	CCACGTAGAG	TACAACGCTC	1140
${\tt TGAGAGAGTT}$	$\mathtt{TCCCGATATA}$	CCTGAGTCAG	TGGAAGATCT	TCGGATGGAC	TCTGAACGTG	1200
TAATTGATCC	${\tt ATATGAATTT}$	${\tt GCTCATGAGA}$	CTATAGACAA	ACTTGAAGAT	GATGTATTTG	1260
AGTAGTGCGC	AAGAGCGTTC	${\tt ATAATTCTGC}$	GTCACGTTAA	AATATCATTA	CAACGTAATC	1320
ACTTTATCGA						1330

<210> 34

<211> 1152

<212> DNA

<213> Yersinia enterocolitica

<400> 34

${\tt GAATTCCCCA}$	${\tt ACTTTGACAC}$	CGATAACCGG	TTCAATAGTA	TCTGGAATAG	ACAGCGAAAG	60
${\tt TTGTTGAAAT}$	AATTGAGTGA	TAGCTTGTTC	AAATGAATAC	ATTTGATCTC	CTAATAGTTA	120
GATAAAATAT	CAACTTAACC	AAAGCACTCT	CGGCAGACCA	TCAATTTTAG	CCTATAATTT	180
${\bf TTAGTTTTTA}$	${\tt TTTTGTCTAA}$	TATAACAACA	AAAACAGCAG	${\tt CGGTTTTTTA}$	TATAACCACC	240
${\tt GGCTATTTTC}$	CCACTAAGAT	AACCTTGTTT	TAATAGCCAA	GGGAATAAAT	AGTCATGAAA	300
ATATCATCAT	${\tt TTATTTCTAC}$	ATCACTGCCC	CTGCCGGCAT	CAGTGTCAGG	ATCTAGCAGC	360
GTAGGAGAAA	${\tt TGTCTGGGCG}$	${\tt CTCAGTCTCA}$	CAGCAAAAA	GTGATCAATA	TGCAAACAAT	420
CTGGCCGGGC	${\tt GCACTGAAAG}$	CCCTCAGGGT	TCCAGCTTAG	CCAGCCGTAT	CATTGAGAGG	480
${\tt TTATCATCAA}$	${\tt TGGCCCACTC}$	${\tt TGTGATTGGA}$	TTTATCCAAC	GCATGTTCTC	GGAGGGGAGC	540
CATAAACCGG	${\tt TGGTGACACC}$	AGCACTCACG	CCTGCACAAA	TGCCAAGCCC	TACGTCTTTC	600
AGTGATAGTA	TCAAGCAACT	TGCTGCTGAG	ACGCTGCCAA	AATACATGCA	GCAGTTGAGT	660

AGCTTGGATG	CAGAGACGCT	${\tt GCAGAAAAAT}$	CATGACCAGT	TCGCCACGGG	CAGCGGCCCT	720
CTTCGTGGCA	GTATCACTCA	ATGCCAAGGG	${\tt CTGATGCAGT}$	${\tt TTTGTGGTGG}$	${\tt GGAATTGCAA}$	780
GCTGAGGCCA	${\tt GTGCCATTTT}$	AAACACGCCT	${\tt GTTTGTGGTA}$	${\tt TTCCCTTCTC}$	GCAGTGGGGA	840
ACTGTTGGTG	GGGCGGCCAG	${\tt CGCGTACGTC}$	GCCAGTGGCG	${\tt TTGATCTAAC}$	GCAGGCAGCA	900
AATGAGATCA	AAGGGCTGGG	${\tt GCAACAGATG}$	${\tt CAGCAATTAC}$	TGTCATTGAT	${\tt GTGATATGGA}$	960
TAAAAACAAG	${\tt GGGGTAGTGT}$	TTCCCCCTTT	TTCTATCAAT	${\tt ATTGCGAATA}$	${\tt TCTTCGTCCC}$	1020
TGATCTTTCA	$\tt GGGGCGAATC$	${\tt GTTTTTTAGC}$	${\tt ATGCTCATTG}$	${\tt TTAGAATTTC}$	${\tt TGACTTATCT}$	1080
CTCTTCTGTA	${\tt TTACTACTCA}$	TACTCTGGAA	AATCCTGAGC	ATTTATATCT	ATGGATTGAT	1140
GCAGCACTCG	AG					1152

<210> 35

<211> 1990

<212> DNA

<213> Yersinia enterocolitica

<400> 35

AGGGCATTGG	AATTAAAAAT	ATATTTATCT	AAATGATGAT	GAGTTTAAAT	TACATTTGCG	60
${\tt TATTAAAATG}$	AATAACGCAT	TATTAACGTA	TTACCATCTG	TTCCCGCTTA	ATTTTTTAAA	120
AAATTTAAGG	${\tt TAACAATGAG}$	${\tt TATATATCTT}$	ATGGGAAAAG	CCAAAAAACT	AACGAACACT	180
ATAATAATTC	GATTAACATC	${\tt AATGAAAATA}$	CACGGCTCAC	СТАТТАТТАА	AATAATACGA	240
CTAGCATTAT	AAGAAAAAAT	${\tt ATTTTTTATG}$	${\tt TTTATAGTAT}$	AGGCGTGTAT	TTAATTAGTT	300
${\tt CTTAATTTAA}$	${\tt TTAAGGAGGG}$	AAGCATGAAC	TTATCATTAA	GCGATCTTCA	TCGTCAGGTA	360
${\tt TCTCGATTGG}$	${\tt TGCAGCAAGA}$	${\tt GAGCGGTGAT}$	TGTACCGGGA	AATTAAGAGG	TAACGTTGCT	420
GCCAATAAAG	AAACTACCTT	${\tt TCAAGGTTTG}$	ACCATAGCCA	GTGGTGCCAG	AGAGTCAGAA	480
AAAGTATTTG	${\tt CTCAAACTGT}$	ACTAAGCCAC	GTAGCAAATA	TTGTTCTAAC	TCAAGAAGAT	540
ACCGCTAAGC	${\tt TATTGCAAAG}$	CACGGTAAAG	CATAATTTGA	ATAATTATGA	ATTAAGAAGT	600
GTCGGCAATG	GTAATAGTGT	ACTTGTCAGT	TTACGTAGTG	ACCAAATGAC	ACTACAAGAC	660

GCCAAAGTGC	TGTTGGAGGC	TGCATTGCGA	CAAGAGTCGG	GAGCGAGGGG	GCATGTATCA	720
TCTCATTCAC	ATTCAGTCCT	TCACGCACCG	GGAACCCCGG	TGCGTGAAGG	ACTGCGTTCA	780
CATCTAGACC	CCAGAACACC	ACCGTTGCCA	CCGCGTGAAC	GACCACACAC	TTCTGGCCAT	840
CACGGGGCTG	GCGAAGCCAG	AGCCACCGCA	CCAAGCACTG	TTTCTCCTTA	TGGCCCAGAA	900
GCGCGCGCAG	AACTCAGCAG	CCGCCTCACC	ACATTGCGCA	ATACGCTGGC	GCCAGCAACG	960
AATGATCCGC	GTTACTTACA	AGCCTGCGGC	GGTGAAAAGC	TAAACCGATT	TAGAGATATT	1020
CAATGCTGTC	GGCAAACCGC	AGTACGCGCC	GATCTTAATG	CCAATTACAT	CCAGGTCGGT	1080
AACACTCGTA	CCATAGCGTG	CCAGTATCCG	CTACAATCTC	AACTTGAAAG	CCATTTCCGT	1140
ATGCTGGCAG	AAAACCGAAC	GCCAGTGTTG	GCTGTTTTAG	CGTCCAGTTC	TGAGATAGCC	1200
AATCAAAGAT	TCGGTATGCC	AGATTATTTC	CGCCAGAGTG	GTACCTATGG	CAGTATCACT	1260
GTAGAGTCTA	AAATGACTCA	GCAAGTTGGT	CTCGGTGACG	GGATTATGGC	AGATATGTAT	1320
ACTTTAACGA	TTCGTGAAGC	GGGTCAAAAA	ACAATTTCTG	TTCCTGTGGT	TCATGTTGGC	1380
AATTGGCCCG	ATCAGACCGC	AGTCAGCTCT	GAAGTTACCA	AGGCACTCGC	TTCACTGGTA	1440
GATCAAACAG	CAGAAACAAA	ACGCAATATG	TATGAAAGCA	AAGGAAGTTC	AGCGGTAGCA	1500
GATGACTCCA	AATTACGGCC	GGTAATACAT	TGCCGTGCGG	GTGTTGGCCG	TACTGCGCAA	1560
CTGATTGGCG	CAATGTGCAT	GAATGATAGT	CGTAATAGTC	AGTTAAGCGT	AGAAGATATG	1620
GTCAGCCAAA	TGCGAGTACA	AAGAAATGGT	ATTATGGTAC	AAAAAGATGA	GCAACTTGAT	1680
GTTCTGATTA	AGTTGGCTGA	AGGACAAGGG	CGACCATTAT	TAAATAGCTA	ATGTAAATAT	1740
TTATTCCTAT	GAGTAAATAA	AATTACTAAG	AGATATACAC	CACTTTGCCA	ATCAAAGAAA	1800
CTTTAAACCT	CAACTAAAGT	AAGCAATTAG	TTGAGGTTTA	TCTGCTATAG	AATAATTATT	1860
AACAAAAATA	TAAACAACAA	AATTAAAAGT	TATGTGTCTA	CTTTTACTTT	ATGTAACCAA	1920
ACCCATTAAT	GGATACCGTA	CGTTTTTCTT	TTATAGAATT	AAACCAGTAA	ATGAGATGAT	1980
GAAGGACGAT	GATCATCGTC			•		1990

<210> 36

<211> 867

<212> DNA

<213> Yersinia enterocolitica

<400> 36

ATGATTGGGC	CAATATCACA	AATAAACAGC	TTCGGTGGCT	TATCAGAAAA	AGAGACCCGT	60
TCTTTAATCA	GTAATGAAGA	GCTTAAAAAT	ATCATAATAC	AGTTGGAAAC	TGATATAGCG	120
GATGGATCCT	GGTTCCATAA	AAATTATTCA	CGCCTGGATA	TAGAAGTCAT	GCCCGCATTA	180
GTAATTCAGG	CGAACAATAA	ATATCCGGAA	ATGAATCTTA	ATTTTGTTAC	ATCTCCCCAG	240
GACCTTTCGA	TAGAAATAAA	AAATGTCATA	GAAAATGGAG	TTGGATCTTC	CCGCTTCATA	300
ATTAACATGG	GGGAGGGTGG	AATACATTTC	AGTGTAATTG	ATTACAAACA	TATAAATGGG	360
AAAACATCTC	TGATATTATT	TGAACCAGTA	AACTTTAATA	GTATGGGGCC	AGCGATACTG	420
GCAATAAGTA	CAAAAACGGC	CATTGAACGT	TATCAATTAC	CTGATTGCCA	TTTTTCCATG	480
${\tt GTGGAAATGG}$	ATATTCAGCG	AAGCTCATCT	GAATGTGGTA	${\tt TTTTTAGTTT}$	GGCACTGGCA	540
AAAAAACTTT	ACACCGAGAG	AGATAGCCTG	${\tt TTGAAAATAC}$	ATGAAGATAA	TATAAAAGGT	600
${\tt ATATTAAGTG}$	${\tt ATAGTGAAAA}$	TCCTTTACCC	CACAATAAGT	TGGATCCGTA	TCTCCCGGTA	660
${\tt ACTTTTTACA}$	AACATACTCA	AGGTAAAAA	${\tt CGTCTTAATG}$	AATATTTAAA	TACTAACCCG	720
${\tt CAGGGAGTTG}$	GTACTGTTGT	$\mathtt{TAACAAAAAA}$	AATGAAACCA	TCTTTAATAG	GTTTGATAAC	780
AATAAATCCA	${\tt TTATAGATGG}$	AAAGGAATTA	${\tt TCAGTTTCGG}$	TACATAAAAA	GAGAATAGCT	840
GAATATAAAA	CACTTCTCAA	AGTATAA				867

<210> 37

<211> 2190

<212> DNA

<213> Yersinia enterocolitica

<400> 37

ATGAAAATCA TGGGAACTAT GCCACCGTCG ATCTCCCTCG CCAAAGCTCA TGAGCGCATC 60
AGCCAACATT GGCAAAATCC TGTCGGTGAG CTCAATATCG GAGGAAAACG GTATAGAATT 120
ATCGATAATC AAGTGCTGCG CTTGAACCCC CACAGTGGTT TTTCTCTCTT TCGAGAAGGG 180

GTTGGTAAGA	TCTTTTCGGG	GAAGATGTTT	AACTTTTCAA	TTGCTCGTAA	CCTTACTGAG	240
ACACTCCATG	CAGCCCAGAA	AACGACTTCG	CAGGAGCTAA	GGTCTGATAT	CCCCAATGCT	300
CTCAGTAATC	TCTTTGGAGC	CAAGCCACAG	ACCGAACTGC	CGCTGGGTTG	GAAAGGGAAG	360
CCTTTGTCAG	GAGCTCCGGA	TCTTGAAGGG	ATGCGAGTGG	CTGAAACCGA	TAAGTTTGCC	420
GAGGGCGAAA	GCCATATTAG	TATAATAGAA	ACTAAGGATA	ATCAGCGGTT	GGTGGCTAAG	480
ATTGAACGCT	CCATTGCCGA	GGGGCATTTG	TTCGCAGAAC	TGGAGGCTTA	TAAACACATC	540
TATAAAACCG	CGGGCAAACA	TCCTAATCTT	GCCAATGTCC	ATGGCATGGC	TGTGGTGCCA	600
TACGGTAACC	GTAAGGAGGA	AGCATTGCTG	ATGGATGAGG	TGGATGGTTG	GCGTTGTTCT	660
GACACACTAA	GAAGCCTCGC	CGATAGCTGG	AAGCAAGGAA	AGATCAATAG	TGAAGCCTAC	720
TGGGGAACGA	TCAAGTTTAT	TGCCCATCGG	CTATTAGATG	TAACCAATCA	CCTTGCCAAG	780
GCAGGGATAG	TACATAACGA	TATCAAACCC	GGTAATGTGG	TATTTGACCG	CGCTAGCGGA	840
GAGCCCGTTG	TCATTGATCT	AGGATTACAC	TCTCGTTCAG	GGGAACAACC	TAAGGGGTTT	900
ACAGAATCCT	TCAAAGCGCC	GGAGCTTGGA	GTAGGAAACC	TAGGCGCATC	AGAAAAGAGC	960
GATGTTTTTC	TCGTAGTTTC	AACCCTTCTA	CATGGTATCG	AAGGTTTTGA	GAAAGATCCG	1020
GAGATAAAGC	CTAATCAAGG	ACTGAGATTC	ATTACCTCAG	AACCAGCGCA	CGTAATGGAT	1080
GAGAATGGTT	ACCCAATCCA	TCGACCTGGT	ATAGCTGGAG	TCGAGACAGC	CTATACACGC	1140
TTCATCACAG	ACATCCTTGG	CGTTTCCGCT	GACTCAAGAC	CTGATTCCAA	CGAAGCCAGA	1200
CTCCACGAGT	TCTTGAGCGA	CGGAACTATT	GACGAGGAGT	CGGCCAAGCA	GATCCTAAAA	1260
GATACTCTAA	CCGGAGAAAT	GAGCCCATTA	TCTACTGATG	TAAGGCGGAT	AACACCCAAG	1320
AAGCTTCGGG	AGCTCTCTGA	TTTGCTTAGG	ACGCATTTGA	GTAGTGCAGC	AACTAAGCAA	1380
TTGGATATGG	GGGTGGTTTT	GTCGGATCTT	GATACCATGT	TGGTGACACT	CGACAAGGCC	1440
GAACGCGAGG	GGGGAGTAGA	CAAGGATCAG	${\tt TTGAAGAGTT}$	TTAACAGTTT	GATTCTGAAG	1500
ACTTACAGCG	TGATTGAAGA	CTATGTCAAA	GGCAGAGAAG	GGGATACCAA	GAGTTCCAGT	1560
GCGGAAGTAT	CCCCTATCA	TCGCAGTAAC	TTTATGCTAT	CGATCGCCGA	ACCTTCACTG	1620
CAGAGGATCC	AAAAGCATCT	GGACCAGACA	CACTCTTTTT	CTGATATCGG	TTCACTAGTG	1680
CGCGCACATA	AGCACCTGGA	AACGCTTTTA	GAGGTCTTAG	TCACCTTGTC	ACCGCAAGGG	1740
CAGCCCGTGT	CCTCTGAAAC	CTACAGCTTC	CTGAATCGAT	TAGCTGAGGC	TAAGGTCACC	1800
TTGTCGCAGC	AATTGGATAC	TCTCCAGCAG	CAGCAGGAGA	GTGCGAAACG	GCAACTATCT	1860
ATTCTGATTA	ATCGTTCAGG	TTCTTGGGCC	GATGTTGCTC	GTCAGTCCCT	GCAGCGTTTT	1920
GACAGTACCC	GGCCTGTAGT	GAAATTCGGC	ACTGAGCAGT	ATACCGCAAT	TCACCGTCAG	1980
ATGATGGCGG	CCCATGCAGC	CATTACGCTA	CAGGAGGTAT	CGGAGTTTAC	TGATGATATG	2040
CGAAACTTTA	CAGCGGACTC	TATTCCACTA	CTGATTCGAC	TTGGACGAAG	CAGTTTAATA	2100
GATGAGCATT	TGGTTGAACA	GAGAGAGAAG	TTGCGAGACG	TGACGACCAT	CGCCGAGCGA	2160
CTGAACCGGT	TGGAGCGGGA	ATGGATGTGA				2190

<210>	38	
<211>	31	
<212>	DNA	
-0125	Yersinia enterocolitica	
<213>	reisinia enterocolitica	
<400>	38	
AAACTO	GCAGA TGTCTCTTGA GCAGAGGAGT C	31
<210>	3.9	
1220		
<211>		
<212>	DNA	
<213>	Yersinia enterocolitica	
-220		
<400>	39	

PCT/IB99/00587

30

<u></u>WO 99/45098

AAACTGCAGT CAGACTCCCT CTTCCTCCTC